



(11) **EP 1 254 666 A1**

(12) **EUROPEAN PATENT APPLICATION**  
published in accordance with Art. 158(3) EPC

(43) Date of publication:  
06.11.2002 Bulletin 2002/45

(51) Int Cl.7: **A61K 39/395**, A61K 9/08,  
A61K 47/04, A61K 47/12

(21) Application number: **00985964.6**

(86) International application number:  
**PCT/JP00/09339**

(22) Date of filing: **27.12.2000**

(87) International publication number:  
**WO 01/047554 (05.07.2001 Gazette 2001/27)**

(84) Designated Contracting States:  
**AT BE CH CY DE DK ES FI FR GB GR IE IT LI LU**  
**MC NL PT SE TR**  
Designated Extension States:  
**AL LT LV MK RO SI**

(30) Priority: **28.12.1999 JP 37520399**

(71) Applicant: **CHUGAI SEIYAKU KABUSHIKI**  
**KAISHA**  
**Tokyo, 115-8543 (JP)**

(72) Inventors:  

- **YAMAZAKI, Tadao,**  
**Chugai Seiyaku Kabushiki Kaisha**  
**Gotenba-shi, Shizuoka 412-8543 (JP)**
- **HAYASAKA, Akira,**  
**Chugai Seiyaku Kabushiki Kaisha**  
**Gotenba-shi, Shizuoka 412-8545 (JP)**
- **KOGA, Akiko, Chugai Seiyaku Kabushiki Kaisha**  
**Gotenba-shi, Shizuoka 412-8543 (JP)**

(74) Representative: **Maschio, Antonio**  
**D Young & Co,**  
**21 New Fetter Lane**  
**London EC4A 1DA (GB)**

(54) **STABLE ANTIBODY COMPOSITIONS AND INJECTION PREPARATIONS**

(57) A stabilized pharmaceutical preparation of an antibody to parathyroid hormone related peptide, wherein the antibody is dissolved in a buffer solution containing at least one buffer selected from the group

consisting of acetic acid, citric acid, phosphoric acid, and salts thereof and is in the form of a solution of pH 5 to 8.

**Description****TECHNICAL FIELD**

5 **[0001]** The present invention relates to a pharmaceutical preparation stabilizing an antibody to parathyroid hormone related peptide and a pharmaceutical preparation for injection.

**BACKGROUND ART**

10 **[0002]** Parathyroid hormone related peptide (hereinafter referred to as PTHrP) is a protein produced by a tumor, which is a major causative agent of hypercalcemia. That protein causes tumor-producing hypercalcemia (Humoral hypercalcemia of malignancy, hereinafter referred to as HHM) by promoting calcium resorption in bone resorption and uridiferous tubule. Although at present, a calcitonin or a bisphosphonate each which has a bone resorption-inhibitory action, is used for HHM therapy, due to rapid progress of HHM and worsening QOL (Quality of Life) of a terminal cancer patient, there is a demand for development of an effective therapeutic agent according with the cause.

15 **[0003]** An antibody to parathyroid hormone related peptide (hereinafter referred to as anti-PTHrP antibody) has an immediate therapeutic effect against HHM after administration thereof, and thus it is excellent in comparison with a bisphosphonate which requires several days until the therapeutic effect is observed. Further, the antibody is useful as a therapeutic agent for cachexia observed in a terminal cancer patient. (Japanese Patent Application Laying-Open  
20 (kokai) No. 11-80025)

**DISCLOSURE OF THE INVENTION**

25 **[0004]** In order to use the anti-PTHrP antibody as a therapeutic agent for diseases, it is necessary to provide it as a stabilized preparation which can retain biological activity of the anti-PTHrP antibody for a long period. Accordingly, it is an object of the present invention to provide a stabilized pharmaceutical preparation of the anti-PTHrP antibody.

**[0005]** The inventors of the present invention prepared an anti-PTHrP antibody solution, verified the influences of hydrogen ion concentration (pH) and buffer solution concentration on physicochemical properties of the anti-PTHrP antibody, and succeeded in producing a stabilized pharmaceutical preparation of the anti-PTHrP antibody.

30 **[0006]** In other words, the present invention provides a stabilized pharmaceutical preparation of an anti-PTHrP antibody, wherein the anti-PTHrP antibody is dissolved in a buffer solution containing at least one buffer selected from the group consisting of acetic acid, citric acid, phosphoric acid, and salts thereof and is in the form of a solution of pH 5 to 8.

35 **[0007]** Further, the present invention provides a stabilized composition of anti-PTHrP antibody solution, wherein the anti-PTHrP antibody is dissolved in a buffer solution containing at least one buffer selected from the group consisting of acetic acid, citric acid, phosphoric acid, and salts thereof and is in the form of a solution of pH 5 to 8.

**[0008]** In detail, the present invention provides the composition of the antibody solution wherein the composition of the antibody solution is a solution composition for bulk.

40 **[0009]** In more detail, the present invention provides the composition of the antibody solution substantially free of stabilizers other than a buffer or an isotonicizing agent.

**[0010]** In this description, "a buffer solution" means a solution having a buffer action (that is, easing up the change of pH), and "a buffer" means a substance having a buffer action.

**[0011]** The preparation or the composition may have a total concentration of the buffer of 0.1 to 100 mmol/L, preferably 5 to 50 mmol/L.

45 **[0012]** To the preparation may be added an isotonicizing agent such as sodium chloride and glucose, so that the preparation essentially has the same osmotic pressure as human blood. In general, a preferable osmotic pressure is approximately from 250 to 350 mOsm.

**[0013]** The anti-PTHrP antibody may be a monoclonal antibody, and this antibody is preferably a human antibody, a humanized antibody, or a chimeric antibody.

50 **[0014]** Moreover, the inventors of the present invention prepared a pharmaceutical preparation for injection containing the anti-PTHrP antibody solution, verified that pain action at a time of administration is different depending on a kind of buffer solution, and succeeded in producing a pharmaceutical preparation for injection which includes the anti-PTHrP antibody and causes less pain.

**[0015]** Namely, the present invention provides a pharmaceutical preparation for injection wherein the anti-PTHrP antibody is dissolved in a buffer solution containing a buffer consisting of acetic acid and/or salts thereof.

55 **[0016]** In detail, the present invention provides the pharmaceutical preparation for injection which is in the form of a solution of pH 5 to 8.

**[0017]** In more detail, the present invention provides the pharmaceutical preparation for injection wherein the solution

has a total concentration of the buffer of 0.1 to 100 mmol/L, preferably 5 to 50 mmol/L.

**[0018]** The anti-PTHrP antibody may be a monoclonal antibody, and this antibody preferably be a human antibody, a humanized antibody, or a chimeric antibody.

**[0019]** Hereinbelow, the present invention will be illustrated in detail.

## 1. Anti-PTHrP antibody

**[0020]** The anti-PTHrP antibody used in the present invention may be any one as far as it has the desired pharmacological effect, regardless of its source, type (monoclonal or polyclonal) and configuration.

**[0021]** The anti-PTHrP antibody used in the present invention can be produced by any known method as a polyclonal or monoclonal antibody. Preferably, the anti-PTHrP antibody is a monoclonal antibody derived from a mammal. The mammal-derived monoclonal antibody includes those produced from a hybridoma and those produced by a genetic engineering technique from a host transformed with a recombinant expression vector carrying a gene for the antibody. The antibody can bind to PTHrP to prevent the binding of the PTHrP to a PTH/PTHrP receptor, thus blocking the signal transduction of the PTHrP and consequently inhibiting the biological activity of the PTHrP.

**[0022]** A specific example of such antibody is #23-57-137-1 antibody which can be produced with a hybridoma clone #23-57-137-1.

**[0023]** The hybridoma clone #23-57-137-1 has been designated "mouse-mouse hybridoma #23-57-137-1" and deposited under the terms of the Budapest Treaty on August 15, 1996 at the National Institute of Bioscience and Human-technology, Agency of Industrial Science and Technology, Japan (1-3, Higashi 1-chome, Tsukuba-shi, Ibaraki, Japan) under the accession No. FERM BP-5631.

## 2. Antibody-producing hybridoma

**[0024]** A monoclonal antibody-producing hybridoma can be produced as follows. That is, PTHrP is used as an antigen for immunization in accordance with a conventional immunization method. The resulting immunocytes are fused to known parent cells by a conventional cell fusion method, and monoclonal antibody-producing cells are screened from the fused cells by a conventional screening method.

**[0025]** First, a human PTHrP, which is used as a sensitizing antigen for producing the antibody, is prepared by expressing the PTHrP gene/amino acid sequence disclosed in Suva, L. J. et al., Science (1987) 237, 893. A nucleotide sequence encoding the PTHrP is inserted into a known expression vector, and a suitable host cell is transformed with the expression vector. The PTHrP protein is then isolated and purified from the transformed host cell or from a culture supernatant of the transformed host cell by any known method.

**[0026]** Then, the purified PTHrP protein is used as a sensitizing antigen. Alternatively, a 34-amino acid peptide of the N-terminal region of the PTHrP may be chemically synthesized as the sensitizing antigen.

**[0027]** The mammal to be immunized with the sensitizing antigen is not particularly limited. However, the mammal is preferably selected taking into consideration of compatibility with the parent cell used for cell fusion. Generally, a rodent (e.g., mouse, rat, hamster), rabbit or monkey may be used.

**[0028]** The immunization of the mammal with the sensitizing antigen can be performed in accordance with any known method, for example, by injecting the sensitizing antigen to a mammal intraperitoneally or subcutaneously. More specifically, the sensitizing antigen is properly diluted with or suspended to phosphate-buffered saline (PBS) or physiological saline, the resulting dilution or suspension is then mixed with an appropriate amount of a conventional adjuvant (e.g., Freund's complete adjuvant) to give an emulsion. The emulsion is injected to a mammal several times at intervals of 4 to 21 days. For the immunization, the sensitizing antigen may be attached to a suitable carrier.

**[0029]** After the immunization, the serum antibody level is checked. When the serum antibody level is confirmed to reach a desired level, immunocytes are isolated from the mammal and then subjected to cell fusion. A preferable immunocyte is a spleen cell.

**[0030]** The parent cell used for the cell fusion (i.e., the counterpart of the cell fusion with the immunocyte) is a myeloma cell derived from a mammal. The myeloma cell is of any known cell line, and, for example, P3 (P3x63Ag8.653) (J. Immunol. (1979) 123, 1548-1550), P3x63Ag8U.1 (Current Topics in Microbiology and Immunology (1978) 81, 1-7), NS-1 (Kohler, G. and Milstein, C. Eur. J. Immunol. (1976) 6, 511-519), MPC-11 (Margulies, D. H. et al., Cell (1976) 8, 405-415), SP2/0 (Shulman, M. et al., Nature (1978) 276, 269-270), FO (de St. Groth, S. F. et al., J. Immunol. Methods (1980) 35, 1-21), S194 (Trowbridge, I. S., J. Exp. Med. (1978) 148, 313-323) or R210 (Galfre, G. et al., Nature (1979) 277, 131-133).

**[0031]** Cell fusion of the immunocyte to the myeloma cell is basically performed in accordance with any known method, such as the method of Milstein et al. (Kohler, G. and Milstein, C., Methods Enzymol. (1981) 73, 3-46).

**[0032]** More specifically, the cell fusion is performed, for example, in a conventional nutrient culture medium in the presence of a cell fusion promoter. The cell fusion promoter may be polyethylene glycol (PEG) or a Sendai virus

(hemagglutinating virus of Japan; HVJ). If desired, for the purpose of improving the fusion efficiency, an additive such as dimethyl sulfoxide may be incorporated.

**[0033]** The ratio between the immunocytes and the myeloma cells for the cell fusion may be any one. For example, the immunocytes are used in the amount 1-10 times larger than the myeloma cells. The culture medium used for the cell fusion is, for example, RPMI 1640 medium or MEM medium suitable for the growth of the above-mentioned myeloma cell lines, or other medium conventionally used for the culture of such cell lines. If desired, a serum supplement, such as fetal calf serum (FCS), may be added to the culture medium.

**[0034]** The cell fusion is performed by fully mixing given amounts of the immunocytes and the myeloma cells in the culture medium, adding a PEG solution (e.g., mean molecular weight: about 1000-6000) (which has been previously warmed to about 37° C) to the mixture usually to a concentration of 30-60% (w/v), and then mixing the resulting solution, thereby producing the desired fusion cells (i.e., hybridomas). Subsequently, an appropriate culture medium is added to the culture solution successively, and centrifuged to remove the supernatant. This procedure is repeated several times to remove the cell fusion promoter or the like that are undesirable for the growth of the hybridomas, from the culture medium.

**[0035]** The obtained hybridomas can be selected by culturing in a conventional selective medium, such as hypoxanthine-aminopterin-thymidine (HAT) medium. The culturing of the hybridomas in HAT medium is performed for the time of period enough to cause the death of the cells other than the desired hybridomas (i.e., cells that fail to fuse), usually for several days to several weeks. Subsequently, conventional limiting dilution method is performed for screening and mono-cloning of the hybridomas that are secreting the desired antibody.

**[0036]** As a method other than preparing the hybridomas by immunizing a non-human mammal with the antigen as described above, a human lymphocyte may be sensitized with PTHrP *in vitro*, and then subjected the sensitized lymphocyte to cell fusion to a human-derived myeloma cell capable of infinite growth, thereby producing a human antibody having a binding activity against the PTHrP (Japanese Patent Publication No. 1-59878). Alternatively, a human antibody against PTHrP may be prepared by injecting PTHrP as an antigen to a transgenic animal that has the entire repertoires of human antibody genes to produce an anti-PTHrP antibody-producing cell, and then immortalizing the cells, thus producing the human antibody from the immortalized cell (International Patent Publication Nos. WO 94/25585, WO 93/12227, WO 92/03918 and WO 94/02602).

**[0037]** The monoclonal antibody-producing hybridoma prepared as above can be subcultured in a conventional culture medium and stored under liquid nitrogen for a long time of period.

**[0038]** For the production of a monoclonal antibody from the hybridoma, a method may be employed that involves culturing the hybridoma in accordance with a conventional technique and collecting the monoclonal antibody from the culture supernatant, or that involves injecting the hybridoma to a mammal compatible with the hybridoma to grow the hybridoma in the mammal and collecting the hybridoma from the ascites of the mammal. The former method is suitable for producing the antibody in high purity, while the latter method is suitable for producing the antibody in a large amount.

### 3. Recombinant antibody

**[0039]** In the present invention, a recombinant-type monoclonal antibody may be used, which can be produced by cloning an antibody gene from the hybridoma, integrating the antibody gene into a suitable vector, introducing the vector into a host, and then producing the antibody from the host according to a conventional genetic recombination technique (see, for example, Vandamme, A. M. et al., *Eur. J. Biochem.* (1990) 192, 767-775, 1990).

**[0040]** Specifically, mRNA encoding variable (V) region of an anti-PTHrP antibody is isolated from the anti-PTHrP antibody-producing hybridoma. The isolation of the mRNA is performed by preparing a total RNA by any known method, such as guanidium ultracentrifugation method (Chirgwin, J. M. et al., *Biochemistry* (1979) 18, 5294-5299) and AGPC method (Chomczynski, P. et al., *Anal. Biochem.* (1987) 162, 156-159), and then producing the desired mRNA from the total RNA using mRNA Purification Kit (Pharmacia) or the like. Alternatively, the mRNA may also be prepared directly using QuickPrep mRNA Purification Kit (Pharmacia).

**[0041]** Next, cDNA for the antibody V-region is synthesized from the obtained mRNA with a reverse transcriptase. The synthesis of the cDNA is performed using AMV Reverse Transcriptase First-strand cDNA Synthesis Kit (Seikagaku Corporation) or the like. The cDNA may also be synthesized and amplified by 5'-RACE method (Frohman, M.A. et al., *Proc. Natl. Acad. Sci. USA* (1988) 85, 8998-9002; Belyavsky, A. et al., *Nucleic Acids Res.* (1989) 17, 2919-2932) using 5'-Ampli FINDER RACE Kit (CLONETECH) in combination with PCR method, or the like.

**[0042]** A DNA fragment of interest is isolated and purified from the resulting PCR product and then ligated to a vector DNA to obtain a recombinant vector. The recombinant vector is introduced into a host such as *E. coli*, and a colony containing a desired recombinant vector is selected. The nucleotide sequence of the DNA of interest in the recombinant vector is confirmed by, for example, dideoxynucleotide chain termination method.

**[0043]** Once DNA encoding the anti-PTHrP antibody V-region is obtained, the DNA is integrated into an expression vector containing a DNA encoding a desired antibody constant (C) region.

**[0044]** For the production of the anti-PTHrP antibody used in the present invention, the antibody gene is integrated into an expression vector so that the antibody gene can be expressed under the control of expression control regions (e.g., enhancer, promoter). A host cell is transformed with the expression vector to express the antibody.

**[0045]** In the expression of the antibody gene, a DNA encoding heavy (H) chain and a DNA encoding light (L) chain of the antibody may be integrated into separate expression vectors, and then a host cell is co-transformed with the resulting recombinant expression vectors. Alternatively, both the DNA encoding H-chain and the DNA encoding L-chain of the antibody may be integrated together into a single expression vector, and then a host cell may be transformed with the resulting recombinant expression vector (WO 94/11523).

**[0046]** For the production of the recombinant antibody, besides the above-mentioned host cells, a transgenic animal may also be used as a host. For example, the antibody gene is inserted into a predetermined site of a gene encoding a protein inherently produced in the milk of an animal (e.g., goat  $\beta$ -casein) to obtain a fusion gene. A DNA fragment containing the antibody gene-introduced fusion gene is injected into an embryo of a goat, and the embryo is then introduced into a female goat. The female goat having the embryo therein bears a transgenic goat. The antibody of interest is secreted in the milk from the transgenic goat or a progeny thereof. For the purpose of increasing the amount of the antibody-containing milk from the transgenic goat, an appropriate hormone may be administered to the transgenic goat (Ebert, K.M. et al., *Bio/Technology* (1994) 12, 699-702).

#### 4. Modified antibody

**[0047]** In the present invention, for the purpose of reducing the heterogeneity against a human body or the like, an artificially modified recombinant antibody may be used, such as a chimeric antibody and a humanized antibody. These modified antibodies can be prepared by the following known methods.

**[0048]** A chimeric antibody usable in the present invention can be prepared by ligating the DNA encoding the antibody V-region prepared as set forth above to a DNA encoding a human antibody C-region, integrating the ligation product into an expression vector, and introducing the resulting recombinant expression vector into a host to produce the chimeric antibody.

**[0049]** A humanized antibody is also referred to as a "reshaped human antibody", in which the complementarity determining regions (CDRs) of an antibody of a non-human mammal (e.g., a mouse) are grafted to those of a human antibody. The general genetic recombination procedures for producing such humanized antibody are also known (EP 125023; WO 96/02576).

**[0050]** Specifically, a DNA sequence in which mouse antibody CDRs are ligated through framework regions (FRs) of a human antibody is amplified by PCR method using several oligonucleotides as primers which have been designed to have regions overlapping to the terminal regions of the CDRs and the FRs. The resulting DNA is ligated to a DNA encoding a human antibody C-region, and the ligation product is integrated into an expression vector. The resulting recombinant expression vector is introduced into a host, thereby producing the humanized antibody (EP 239044, WO 96/02576).

**[0051]** The FRs of the human antibody ligated through the CDRs are selected so that the CDRs can form a suitable antigen binding site. If necessary, an amino acid(s) in the FRs of the antibody V-region may be replaced so that the CDRs of the reshaped human antibody can form a suitable antigen binding site (Sato, K. et al., *Cancer Res.* (1993) 53, 851-856).

**[0052]** The C-region of the chimeric or humanized antibody may be any human antibody C-region, such as  $C\gamma 1$ ,  $C\gamma 2$ ,  $C\gamma 3$  or  $C\gamma 4$  for the H-chain, and  $C\kappa$  or  $C\lambda$  for the L-chain. The human antibody C-region may be modified for the purpose of improving the stable production of the antibody.

**[0053]** The chimeric antibody is composed of V-regions derived from a non-human mammalian antibody and C-regions derived from a human antibody. The humanized antibody is composed of CDRs derived from a non-human mammalian antibody and FRs and C-regions derived from a human antibody. The humanized antibody is useful as an active ingredient for the drug of the present invention, because the antigenicity of the antibody against a human body is reduced.

**[0054]** A specific example of the humanized antibody usable in the present invention is humanized #23-57-137-1 antibody; in which the CDRs are derived from mouse-derived #23-57-137-1 antibody; the L-chain is composed of the CDRs ligated through three FRs (FR1, FR2 and FR3) derived from human antibody HSU 03868 (GEN-BANK, Deftos, M. et al., *Scand. J. Immunol.*, 39, 95-103, 1994) and a FR (PR4) derived from human antibody S25755 (NBRF-PDB); and the H-chain is composed of the CDRs ligated through FRs derived from human antibody S31679 (NBRF-PDB, Cuisinier, A. M. et al., *Eur. J. Immunol.* 23, 110-118, 1993) in which a part of the amino acid residues in the FRs is replaced so that the reshaped humanized antibody can exhibit an antigen-binding activity.

**[0055]** The *E. coli* strains containing the plasmids having DNAs encoding the H-chain and the L-chain of the humanized #23-57-137-1 antibody, respectively, are designated *Escherichia coli* JM109 (hMBC1HcDNA/pUC19) (for H-chain) and *Escherichia coli* JM109 (hMBC1Lq/pUC19) (for L-chain), respectively. These strains have been deposited under

the terms of the Budapest Treaty on August 15, 1996 at the National Institute of Bioscience and Human-technology, Agency of Industrial Science and Technology, Japan (1-3, Higashi 1-chome, Tsukuba-shi, Ibaraki, Japan), under the accession No. FERM BP-5629 for *Escherichia coli* JM109 (hMBC1HcDNA/pUC19), and under the accession No. FERM BP-5630 for *Escherichia coli* JM109 (hMBC1Lqλ/pUC19).

## 5. Antibody variants

**[0056]** The antibody used in the present invention may be a fragment thereof or a modified form of the fragment, as long as it can bind to PTHrP and inhibit the activity of the PTHrP. For example, the fragment of the antibody includes Fab, F(ab')<sub>2</sub>, Fv, or a single chain Fv (scFv) composed of a H-chain Fv fragment and a L-chain Fv fragment linked together through a suitable linker. Specifically, such antibody fragments can be produced by cleaving the antibody with an enzyme (e.g., papain, pepsin) into antibody fragments, or by constructing a gene encoding the antibody fragment and inserting the gene into an expression vector and introducing the resulting recombinant expression vector into a suitable host cell, thereby expressing the antibody fragment (see, for example, Co, M. S., et al., *J. Immunol.* (1994), 152, 2968-2976; Better, M. & Horwitz, A. H., *Methods in Enzymology* (1989), 178, 476-496, Academic Press, Inc.; Plueckthun, A. & Skerra, A., *Methods in Enzymology* (1989) 178, 476-496, Academic Press, Inc.; Lamoyi, E., *Methods in Enzymology* (1989) 121, 652-663; Rousseaux, J. et al., *Methods in Enzymology* (1989) 121, 663-669; and Bird, R. E. et al., *TIBTECH* (1991) 9, 132-137).

**[0057]** The scFv can be produced by linking the H-chain V-region to the L-chain V-region through a linker, preferably a peptide linker (Huston, J. S. et al., *Proc. Natl. Acad. Sci. USA* (1988) 85, 5879-5883). The H-chain V-region and the L-chain V-region in the scFv may be derived from any one of the antibodies described herein. The peptide linker which binds the V-regions may be any single chain peptide, for example, of 12-19 amino acid residues.

**[0058]** The DNA encoding the scFv can be prepared by first amplifying a DNA encoding the H-chain V-region and a DNA encoding the L-chain V-region of the antibody separately using a DNA fragment encoding the entire region or a part of the H-chain that includes the V-region and a DNA fragment encoding the entire region or a part of the L-chain that includes the V-region as templates and primer pairs that define the terminal ends of the DNA fragments; and then amplifying a DNA encoding the peptide linker using a DNA fragment encoding the peptide linker as a template and a primer pair that define the terminal ends of the DNA fragment so that each terminal end of the peptide linker is ligated to the H-chain V-region and the L-chain V-region, respectively.

**[0059]** Once the DNA encoding the scFv is prepared, an expression vector carrying the DNA and a host transformed with the expression vector can be prepared by conventional methods. The scFv can be produced from the transformed host by a conventional method.

**[0060]** The fragments of the antibody may be produced by preparing genes for the fragments and expressing the genes in suitable hosts as described above. The antibody fragments is also encompassed in the "antibody" of the present invention.

**[0061]** As a modified form of the above-mentioned antibodies, for example, anti-PTHrP antibody conjugated to any molecule (e.g., polyethylene glycol) may also be used. Such modified antibodies are also encompassed in the "antibody" of the present invention. The modified antibodies can be prepared by chemical modifications of the antibodies. The chemical modification techniques suitable for this purpose have already been established in the art.

## 6. Expression and production of recombinant antibody or modified antibody

**[0062]** The antibody gene constructed as described above can be produced and expressed by known methods. For the expression in a mammalian cell, a conventional useful promoter, the antibody gene to be expressed and a poly(A) signal (located downstream to the 3' end of the antibody gene) are operably linked. For example, as the useful promoter/enhancer system, a human cytomegalovirus immediate early promoter/enhancer system may be used.

**[0063]** Other promoter/enhancer systems usable in the expression of the antibody used in the present invention include those derived from viruses (e.g., retrovirus, polyoma virus, adenovirus and simian virus 40 (SV40)) and those derived from mammalian cells (e.g., human elongation factor 1  $\alpha$  (HEF1  $\alpha$ )).

**[0064]** When SV40 promoter/enhancer system is used, the gene expression may be performed readily by the method of Mulligan et al. (*Nature* (1979) 277, 108). When HEF1 a promoter/enhancer system is used, the gene expression may be performed readily by the method of Mizushima et al. (*Nucleic Acids Res.* (1990) 18, 5322).

**[0065]** For the expression in *E. coli*, a conventional useful promoter, a signal sequence for secreting the antibody of interest and the antibody gene may be operably linked. As such a promoter, lacZ promoter or araB promoter may be used. When lacZ promoter is used, the gene expression may be performed by the method of Ward et al. (*Nature* (1998) 341, 544-546; *FASBE J.* (1992) 6, 2422-2427). When araB promoter is used, the gene expression may be performed by the method of Better et al. (Better et al., *Science* (1988) 240, 1041-1043).

**[0066]** Regarding the signal sequence for secretion of the antibody, when the antibody of interest is intended to be

secreted in a periplasmic space of the *E. coli*, *pelB* signal sequence (Lei, S. P. et al., *J. Bacteriol.* (1987) 169, 4379) may be used. The antibody secreted into the periplasmic space is isolated and then refolded so that the antibody takes an appropriate configuration for use.

**[0067]** Regarding the replication origin, those derived from viruses (e.g., SV40, polyoma virus, adenovirus, bovine papilloma virus (BPV)) or the like may be used. In order to increase the gene copy number in the host cell system, the expression vector may further contain a selective marker gene, such as an aminoglycoside phosphotransferase (APH) gene, a thymidine kinase (TK) gene, an *E. coli* xanthine-guanine phosphoribosyltransferase (*Ecogpt*) gene and a dihydrofolate reductase (*dhfr*) gene.

**[0068]** For the production of the antibody used in the present invention, any expression system such as eukaryotic and prokaryotic cell systems may be used. The eukaryotic cell includes established cell lines of animals (e.g., mammals, insects, molds and fungi, yeast). The prokaryotic cell includes bacterial cells such as *E. coli* cells.

**[0069]** It is preferable that the antibody used in the present invention be expressed in a mammalian cell, such as a CHO, COS, myeloma, BHK, Vero or HeLa cell.

**[0070]** Next, the transformed host cell is cultured *in vitro* or *in vivo* to produce the antibody of interest. The culturing of the host cell may be performed by any known method. The culture medium usable herein may be DMEM, MEM, RPMI 1640 or IMDM medium. The culture medium may contain a serum supplement, such as fetal calf serum (FCS).

#### 7. Isolation and purification of antibody

**[0071]** The antibody expressed and produced as described above may be isolated from the cells or the host animal body and purified to uniformity. The isolation and purification of the antibody used in the present invention may be performed on an affinity column. Examples of a protein A column include Hyper D, POROS and Sepharose F.F. (Pharmacia). The method is not particularly limited and other methods conventionally used for the isolation and purification of an antibody may also be employed. For example, various chromatographs using columns other than the above-mentioned affinity column, filtration, ultrafiltration, salting out and dialysis may be used singly or in combination to isolate and purify the antibody of interest (Antibodies A Laboratory Manual. Ed. Harlow, David Lane, Cold Spring Harbor Laboratory, 1988).

#### 8. Determination of the activities of the antibody

**[0072]** The determination of the antigen-binding activity (Antibodies A Laboratory Manual, Ed. Harlow, David Lane, Cold Spring Harbor Laboratory, 1988) or the inhibitory activity against a ligand receptor (Harada, A. et al., *International Immunology* (1993) 5, 681-690) of the antibody used in the present invention may be performed by any known methods.

**[0073]** The method for the determination of the antigen-binding activity of the anti-PTHrP antibody used in the present invention may be BIAcore method (analytical method using surface plasmon resonance), ELISA (enzyme-linked immunosorbent assay), EIA (enzyme immunoassay), RIA (radioimmunoassay) or a fluorescent antibody. For example, when enzyme immunoassay is employed, a sample solution containing the anti-PTHrP antibody (e.g., a culture supernatant of anti-PTHrP antibody-producing cells, or the anti-PTHrP antibody in a purified form) is added to a plate on which PTHrP (1-34) is previously coated. A secondary antibody labeled with an enzyme (e.g., alkaline phosphatase) is further added to the plate. The plate is incubated and washed. A substrate for the enzyme (e.g., p-nitrophenylphosphoric acid) is added to the plate, and the absorbance of the solution in the plate is measured to evaluate the antigen-binding activity of the antibody.

**[0074]** To confirm the activity of the antibody used in the present invention, a neutralizing activity of the antibody (e.g., anti-PTHrP antibody) may be determined.

#### 9. Bulk material

**[0075]** The term "bulk material" refers to a composition containing an anti-PTHrP antibody and an anti-PTHrP antibody composition obtained by isolating and purifying, from a cell or a host animal, the antibody expressed and produced by the above method. The bulk material obtained by isolation and purification includes a solvent such as a buffer solution, used during the isolation and the purification. Further, a metal halide or the like, such as sodium chloride, potassium chloride, and calcium chloride, preferably sodium chloride, is added as an isotonicizing agent to the purified anti-PTHrP antibody solution composition of the present invention so that the composition essentially has the same osmotic pressure as human blood. In general, a preferable osmotic pressure is about 250 to 350 mOsm.

**[0076]** Additionally, at least one kind of the buffer selected from the group consisting of acetic acid, citric acid, phosphoric acid, and salts thereof is added so that the concentration becomes 0.1 to 100 mmol/L, preferably about 5 to 50 mmol/L, and the pH is adjusted to 5 to 8, preferably 5.5 to 7.0, more preferably about 6.0, thereby preparing the antibody solution composition for bulk.

**[0077]** The antibody solution composition for bulk is stored in the state of solution or is frozen until it is used as pharmaceutical preparation, preferably in a frozen state.

**[0078]** The composition contains about 1 to 100 mg/mL of the antibody, and further may contain a surfactant (e.g., Polysorbate 20, Polysorbate 80, Triton, sodium dodecyl sulfate, sodium octyl glycoside, lauryl-, linoleyl-, or stearyl-sulfobetaine, lauryl-, myristyl-, linoleyl- or stearyl-sarcosine, myristyl-, linoleyl-, or cetyl-betaine, lauroamidopropyl-, cocamidopropyl-, linoleamidopropyl-, myristamidopropyl-, palmidopropyl-, or isostearamidopropyl-betaine, myristamidopropyl-, palmidopropyl-, or isostearamidopropyl-dimethylamine, sodium methyl cocoyl- or disodium methyl oleyl-taurate, polyethylene glycol, polypropylene glycol, and a copolymer of ethylene glycol or propylene glycol (Pluronic etc.)), sugar or a sugar alcohol (such as polyols of sucrose, trehalose, glycerol, arabitol, xylitol, sorbitol, and mannitol), an amino acid such as glutamic acid and histidine, or the like which becomes a cryoprotectant for reducing fine particle formation of the antibody during freeze-thawing or which becomes a lyoprotectant during freeze-drying. The concentration thereof may depend on the concentration of the antibody and isotonicity of the pharmaceutical preparation.

**[0079]** It is preferred that the antibody solution composition for bulk, of the present invention is stable at least for 2 years at 2 to 8 °C.

**[0080]** The solution composition for bulk is required to have a long-term preservability and stability against physical stresses during transportation, and simultaneously, in a certain embodiment, it is desirable not to add as little an additive, such as a stabilizer, as possible so as to prepare a pharmaceutical preparation for an administration method, for example subcutaneous administration, suitable for a patient to be treated by using this composition.

**[0081]** Accordingly, it is preferred that the solution composition for bulk of the present invention does not contain any stabilizer except a buffer, an isotonicizing agent such as a metal halide, a surfactant, sugar, or sugar alcohol. Further, it is preferred that the solution composition for bulk of the present invention does not contain any stabilizer except a buffer, an isotonicizing agent such as a metal halide, a surfactant. It is most preferred that the solution composition for bulk of the present invention does not contain any stabilizer except a buffer and an isotonicizing agent such as a metal halide.

#### 10. Administration method and preparation

**[0082]** The anti-PTHrP antibody can be used as an active ingredient for the following: a therapeutic agent for diseases caused by PTH or PTHrP (for example hypercalcemia, hypercalcemia crisis, drug-resistant hypercalcemia, cachexia, a symptom caused by low vasopressin concentration); a QOL improving agent for alleviating a symptom of a disease caused by PTH or PTHrP; an alleviation agent for a central nervous system disease caused by PTH or PTHrP; an alleviation agent for a disease caused by PTH or PTHrP- cytokine cascade; a central nervous system regulator; a cytokine network regulator. The anti-PTHrP antibody can be administered as a purpose for any one of or a plurality of the above applications.

**[0083]** A drug containing the anti-PTHrP antibody as an active ingredient may be administered orally or parenterally, preferably parenterally. The drug may take any dosage form, such as a transpulmonary agent (e.g., an agent administered with the help of a device such as a nebulizer), a nasogastric agent, a transdermic agent (e.g., ointment, cream) or an injection. Examples of the injection include an intravenous injection (e.g., drops), an intramuscular injection, an intraperitoneal injection and a subcutaneous injection for systemic or topical administration.

**[0084]** In particular, in the case of the preparation for injection, acetic acid and/or a salt thereof may preferably be used as a buffer. When acetic acid and/or a salt thereof is used as a buffer, a patient feels less pain at a time of administration of the injection preparation.

**[0085]** Further, depending on the age or the condition of a patient, the route of administration may be selected as appropriate. An effective single dose may be selected within the range from 0.001 to 1000 mg per kg of the body weight. Alternatively, the dose per patient may be selected in the range from 0.01 to 100000 mg/body, preferably 0.1 to 10000 mg/body, more preferably 0.5 to 1000 mg/body, much more preferably 1 to 100 mg/body. However, the drug containing the anti-PTHrP antibody of the present invention is not particularly limited to these ranges.

**[0086]** With respect to the timing of the administration, the drug may be administered to a patient at any stage, including before or after the occurrence of the disease or symptom. The drug may also be administered at the stage where the development of weight loss is predicted in the patient.

**[0087]** The drug comprising the anti-PTHrP antibody as an active ingredient of the present invention may be formulated by any conventional method (Remington's Pharmaceutical Science, latest edition, Mark Publishing Company, Easton, USA). The preparation may further comprise pharmaceutically acceptable carriers and additives.

**[0088]** Examples of such carriers and additives include water, pharmaceutically acceptable organic solvents, amino acids, collagen, polyvinyl alcohol, polyvinylpyrrolidone, carboxyvinyl polymer, sodium carboxymethyl cellulose, sodium polyacrylate, alginate sodium, water soluble dextran, carboxymethyl starch sodium, pectin, methylcellulose, ethylcellulose, xanthan gum, gum Arabic, casein, agar, polyethylene glycol, diglycerine, glycerine, propylene glycol, Vaseline, paraffin, stearyl alcohol, stearic acid, human serum albumin (HSA), mannitol, sorbitol, lactose, and surfactants accept-



able as pharmaceutical additives.

**[0089]** In the practical use, the additive is properly selected from the above members either singly or in combination depending on (without limitation) the dosage form employed. For example, for use as an injectable form, the anti-PTHrP antibody of the purified form is dissolved in a solvent (e.g., physiological saline, a buffer, a grape sugar solution) and then an adsorption-preventing agent (e.g., Polysorbate 80, Polysorbate 20, a gelatin, human serum albumin) is added thereto. The therapeutic agent of the present invention may also be in a re-constitutable freeze-dried form, which is dissolved before use. For the formulation of the freeze-dried dosage form, an excipient such as a sugar alcohol (e.g., mannitol, grape sugar) or a sugar may be incorporated,

**[0090]** In order to provide a stabilized pharmaceutical preparation of the anti-PTHrP antibody, the anti-PTHrP antibody is dissolved in the buffer solution so that the solution has a pH of 5 to 8. The buffer solution may be a mixture solution of an acid (preferably a weak acid such as acetic acid, citric acid, and phosphoric acid) and a salt thereof (preferably an alkali salt such as sodium salt, and potassium salt). The total concentration of the buffer (for example, an acid and a salt thereof) in the preparation may be from 0.1 to 100 mmol/L, preferably 5 to 50 mmol/L, more preferably 10 to 20 mmol/L. The buffer solution of acetic acid, citric acid, or phosphoric acid is prepared by a generally known method (D. D. Perrin, B. Dempsey, "Selection and Application of Buffer Solution" Kodansha Scientific).

**[0091]** An example of prescription for stabilized pharmaceutical preparation of the anti-PTHrP antibody is shown as follows.

anti-PTHrP antibody	20~100 mg
buffer *	5~50 mmol/L
sodium chloride	130~150 mmol/L
total	1~5 mL (pH 5 to 8)

\*: acetic acid buffer solution, citric acid buffer solution, phosphoric acid buffer solution, or the combination thereof

**[0092]** Incidentally, this description incorporates all the contents of the specification and/or drawings of Japanese Patent Application No. 11-375203, from which the priority is claimed.

#### BRIEF DESCRIPTION OF THE DRAWINGS

**[0093]**

Fig. 1 is an electropherogram showing SDS-PAGE patterns before and after acceleration test with respect to preparations at various pH values.

Fig. 2 is an electropherogram showing SDS-PAGE patterns before and after acceleration test with respect to preparations with various concentrations of buffer solutions.

#### BEST MODE FOR CARRYING OUT THE INVENTION

**[0094]** Hereinbelow, the present invention will be described in greater detail with reference to the following Reference Examples and Examples, which should not be construed as limiting the technical scope of the invention.

[EXAMPLE 1] Effect of hydrogen ion concentration (pH)

**[0095]** The anti-PTHrP antibodies used in Examples 1 and 2 were humanized antibodies (hereinafter referred to as "humanized antibody") prepared in accordance with Reference Examples 1 to 4 described below. In addition, analytical methods and analytical conditions for Examples 1 and 2 were as follows.

GPC-UV

**[0096]**

Stationary phase: 50 mmol/L of phosphoric acid buffer solution containing 300 mmol/L of NaCl (pH 6.8)

Flow rate: 0.5 mL/min

Column: G-3000SWXL

Detection: 280 nm

Sample injection volume: 90 µg (inject 30 µl of 3 mg/ml)

## EP 1 254 666 A1

BIACORE method (analytical method using surface plasmon resonance. BIACORE manufactured by Pharmacia Biotech K. K.)

### [0097]

#### 1) Reagent

NHS (N-hydroxysuccinic acid imido), EDC (N-ethyl-N'(3-dimethylamino propyl)-carbodiimido hydrochloride), Ethanolamine: amine coupling kit (Biacore Co.)

PDEA (2-(2-pyridinyldithio)ethaneamine hydrochloride): thiol coupling kit (Biacore Co.), PTHrP (1-34+C): synthesized product (Sawady technology Co., Ltd.)

#### 2) Sensor chip ligand: antigen PTHrP or antigen Protein A

#### 3) Preparation of sample solution

i) A sample solution was diluted with distilled water for injection to about 100 µg/mL, and the concentration was determined by absorbance method at 280 nm.

ii) preparation of unknown sample: with reference to the concentration determined by the absorbance method, the unknown sample was diluted with HBS-EP buffer (Biacore Co., Code #BR-1001-88) identical to one used at a time of BIACORE measurement and 20 µg/mL solution was prepared. All the HBS-EP buffers used for dilution were the same as those used at a time of BIACORE measurement. Then, 20 µL of this solution and 180 µL of HBS-EP buffer were mixed and thus obtained 2 µg/mL solution was regarded as a unknown sample.

iii) preparation of samples for calibration curve: about 100 µg/mL solution, as a standard, was diluted to make not less than 5 different concentrations thereof.

### [0098] Ion Exchange Chromatography (hereinafter referred to as IEC-UV)

Column: PolyCAT A 4.6 × 250 mm

Flow rate: 1.0 ml/min

Detection wavelength: 280 nm

Injection volume: about 30 µg

Elution conditions:

Solvent A: 50 mmol/L MES-NaOH (pH 6.1)

Solvent B: 50 mmol/L MES-NaOH (pH 6.1), 500mmol/L NaCl

SDS-PAGE (reduction, nonreduction / CBB, WB)

[0099] The sample and the pre-treatment solution were mixed with a ratio of 1:1, and heated at 100 °C for 1 minute. Then, SDS-PAGE (Gradient Gel 10-15 was used) was performed. A sample application volume was adjusted so that the concentration became about 1 to 2 mg/ml. Staining and bleaching were conducted. The sample was dipped in a bleaching solution containing 5 % of glycerine for not less than 30 minutes, and dried.

[0100] Pre-treatment solution: 40 mmol/L Tris-HCl buffer solution (pH 8.0) containing 5 % SDS (for reduction treatment, 10 % 2-mercaptoethanol was contained)

Staining: 0.1 % CBB (PhastGel Blue R)

Molecular weight marker: SDS-PAGE Standards Broad Range (BIO-RAD/Cat.No.161-0317)

### [0101]

Protein	Mol.Wt.
Myosin, Rabbit Muscle	200,000
Galactosidase, <i>E.coli</i>	116,250
Phosphorylase b, Rabbit Muscle	97,400
Albumin, Bovine Serum	66,200
Ovalbumin, Chicken Egg	45,000
Carbonic Anhydrase, Bovine Erythrocytes	31,000
Trypsin Inhibitor, Soybean	21,500

# EP 1 254 666 A1

(continued)

Protein	Mol.Wt.
Lysozyme , Chicken Egg	14,400
Aprotinin, Bovine Lung	6,500

(1) Phosphoric acid buffer solution and citric acid buffer solution

**[0102]** A pharmaceutical preparation having the following composition was prepared and an acceleration test was conducted.

1mg/mL of humanized antibody

100 mmol/L of citric acid/sodium citrate buffer solution (pH 4 to 5.5) or 100 mmol/L of sodium phosphate buffer solution (pH 6 to 8)

**[0103]** The conditions for the acceleration test were as follows. The pharmaceutical preparation was stored at 50 °C for from 1 week to 1 month, and during that period the stability was checked. In general, for storage of the preparation at 2 to 8 °C, when the stability can be retained at least at 25 °C for 6 months, at 30 °C for 1 month, or at 40 °C for 1 month, the stability should be retained at 2 to 8 °C for 2 years. Furthermore, for storage of the preparation at 25 or 30 °C, generally when the stability can be retained at 40 °C for 6 months, the stability should be retained at 25 °C for 2 years or at 30 °C for 2 years.

**[0104]** In Table 1, with respect to the preparation before and after the acceleration test, there are described survival rate (%) of the humanized antibodies by GPC-UV, remaining bioactivity rate (%) of the humanized antibodies by BI-ACORE, and main peak survival rate (%) of the humanized antibody by IEC-UV.

TABLE 1  
(Effect of hydrogen ion concentration (pH))

Conc. of hydrogen ion (pH)	pH 4		pH 5		pH 5.5		pH 6		pH 6.5		pH 7		pH 8	
	citric acid buffer solution	citric acid buffer solution	citric acid buffer solution	citric acid buffer solution	citric acid buffer solution	citric acid buffer solution	phosphoric acid buffer solution	phosphoric acid buffer solution	phosphoric acid buffer solution	phosphoric acid buffer solution	phosphoric acid buffer solution	phosphoric acid buffer solution	phosphoric acid buffer solution	phosphoric acid buffer solution
survival rate (%) of the humanized antibodies by GPC-UV	before accel. Test	100	100	100	100	100	100	100	100	100	100	100	100	100
	50 °C-1 week	0	90.1	98.3	98.3	99.3	99.3	97.1	86.2	87.2				
remaining bioactivity rate (%) of the humanized antibodies by BIACORE	before accel. test	-	100	-	-	100	100	-	100	100				
	50 °C-1 week	-	86.7	-	-	110.7	110.7	-	98	86.7				
main peak survival rate (%) of the humanized antibody by IEC-UV	before accel. test	-	100	-	-	100	100	-	100	100				
	25 °C-3 months	-	59.5	-	-	65.9	65.9	-	54.6	53.3				

(2) Acetic acid buffer solution

[0105] A pharmaceutical preparation having the following composition was prepared and the acceleration test was conducted.

13 mg/mL of humanized antibody  
20 mmol/L of acetic acid/sodium acetate buffer solution

150 mmol/L of sodium chloride

[0106] In Table 2, with respect to the preparation before and after the acceleration test, there are described hydrogen ion concentration,  $UV_{360nm}$ , survival rate (%) of the humanized antibodies by GPC-UV, remaining bioactivity rate (%) of the humanized antibodies by BIACORE, and main peak survival rate (%) of the humanized antibody by IEC-UV.

TABLE 2

(Effect of hydrogen ion concentration (pH))						
Samples		01L	02L	03L	04L	05L
hydrogen ion concentration (pH)	before accel. Test	4.94	5.40	6.01	6.52	6.92
	50 °C - 1 month	4.99	5.44	6.04	6.55	6.92
$UV_{360nm}$	50 °C - month	1.4458	0.1730	0.1038	0.0878	0.0821
survival rate (%) of the humanized antibodies by GPC-UV	before accel. Test	100	100	100	100	100
	25 °C - 1 month	96.8	97.0	98.6	98.5	96.6
	25 °C - 3 months	93.7	97.7	97.3	101.8	100.6
	50 °C - 1 month	-	82.5	91.4	92.2	90.8
remaining bioactivity rate (%) of the humanized antibodies by BIACORE	before accel. test	PTHrP	100	100	100	100
		Protein A	100	100	100	100
	50 °C - 1 month	PTHrP	-	78.1	90.0	90.6
		Protein A	-	79.1	94.0	94.8
main peak survival rate (%) of the humanized antibody by IEC-UV	before accel. Test	100	100	100	100	100
	25 °C - 1 month	103.4	85.6	100.9	96.6	107.5
	25 °C - 1 month	84.9	71.6	95.3	102.1	118.0

[0107] In addition, SDS-PAGE patterns of the humanized antibodies before and after acceleration test are described in Fig. 1. In Fig. 1, the upper column shows results under reduction condition, and the lower column shows results under nonreduction condition. The left column shows patterns of the preparation before acceleration test, the center column shows patterns of the preparation after acceleration test with the condition at 50 °C for 1 week, and the right column shows patterns of the preparation after acceleration test with the condition at 50 °C for 1 month. Furthermore, lanes from the left show a pattern of the preparation at pH of 5.0, 5.5, 6.0, 6.5, and 7.0, respectively.

[0108] According to the analytical results of  $UV_{360nm}$ , GPC-UV, BIACORE, and IEC-UV, the humanized antibody was stable at pH of 6 to 7, and most stable at pH 6.

[0109] In view of the analytical results of SDS-PAGE, it was recognized that band on H-chain was increased at pH of 7 and 6.5, and that low molecular decomposition products were increased at pH of 5 and 5.5, and thus it was decided that an optimum pH value was 6.

[Example 2] Effect of concentration of buffer solution

(1) Acetic acid buffer solution

[0110] A pharmaceutical preparation having the following composition was prepared and an acceleration test was conducted.

6.5 mg/mL of humanized antibody  
10, 50, and 100 mmol/L of acetic acid/sodium acetate buffer solution (pH 6)  
150 mmol/L of sodium chloride

[0111] In Table 3, with respect to the preparation before and after the acceleration test, the following are described: hydrogen ion concentration,  $UV_{360nm}$ , survival rate (%) of the humanized antibodies by GPC-UV, and remaining bioactivity rate (%) of the humanized antibodies by BIACORE.

Table 3

(Effect of concentration of acetic acid buffer solution)					
Samples		06L	07L	08L	
Concentration of buffer solution. [mmol/L]		10	50	100	
hydrogen ion concentration (pH)	before accel. Test	6.00	6.00	5.99	
	50 °C - 1 month	6.04	6.01	6.00	
UV <sub>360nm</sub>	50 °C - 1 month	0.0401	0.0359	0.0353	
survival rate (%) of the humanized antibodies by GPC-UV	before accel. test	100	100	100	
	25 °C - 1 month	99.1	96.7	95.9	
	25 °C - 3 months	95.8	97.1	96.7	
	50 °C - 1 month	92.5	92.2	92.5	
remaining bioactivity rate (%) of the humanized antibodies by BIACORE	before accel. test	PTHrP	100	100	100
		Protein A	100	100	100
	50 °C - 1 month	PTHrP	90.4	90.2	90.4
		Protein A	94.5	94.0	92.0

## (2) Citric acid buffer solution

**[0112]** A pharmaceutical preparation having the following composition was prepared and the acceleration test was conducted.

8 mg/mL of humanized antibody

10, 50, and 100 mmol/L of citric acid/sodium citrate buffer solution (pH 6)

150 mmol/L of sodium chloride

**[0113]** In Table 4 with respect to the preparation before and after the acceleration test, the following are described: hydrogen ion concentration, UV<sub>360nm</sub>, survival rate (%) of the humanized antibodies by GPC-UV, and remaining bioactivity rate (%) of the humanized antibodies by BIACORE.

Table 4

(Effect of concentration of citric acid buffer solution)					
Samples		09L	10L	11L	
Concentration of buffer solution. [mmol/L]		10	50	100	
hydrogen ion concentration (pH)	before accel. test	6.10	6.01	6.03	
	50 °C - 1 month	6.12	6.03	6.04	
UV <sub>360nm</sub>	50 °C - 1 month	0.0558	0.0597	0.0536	
survival rate (%) of the humanized antibodies by GPC-UV	before accel. test	100	100	100	
	25 °C - 1 month	96.2	97.1	96.0	
	25 °C - 3 months	97.0	95.9	96.9	
	50 °C - 1 month	92.5	91.0	92.5	
remaining bioactivity rate (%) of the humanized antibodies by BIACORE	before accel. test	PTHrP	100	100	100
		Protein A	100	100	100
	50 °C - 1 month	PTHrP	90.3	88.4	89.7
		Protein A	93.6	90.0	92.1

[0114] In addition, SDS-PAGE patterns of the humanized antibodies before and after acceleration test are described in Fig. 2. In Fig. 2, the upper column shows results under reduction condition, and the lower column shows results under nonreduction condition. The left column shows patterns of the preparation before acceleration test, the center column shows patterns of the preparation after acceleration test with the condition at 50 °C for 1 week, and the right column shows patterns of the preparation after acceleration test with the condition at 50 °C for 1 month. Furthermore, lanes from the left show a pattern of the preparation by using acetic acid buffer solution having a concentration of 10, 50, and 100 mmol/L and citric acid buffer solution having a concentration of 10, 50, 100 mmol/L, respectively.

[0115] According to the analytical results of BIACORE, the humanized antibody was stable by 10 to 50 mmol/L of acetic acid buffer solution and citric acid buffer solution, more stable by 10 mmol/L of the buffer solutions thereof.

[0116] In view of the analytical results of SDS-PAGE, the amount of decomposition products was smaller by citric acid rather than acetic acid, and with low concentration rather than high concentration.

[Example 3]

[0117] In Example 3, a pharmaceutical preparation for injection was prepared, and the preparation was reviewed concerning pain-easing effect. Incidentally, the anti-PTHrP antibody used in this Example 3 was a humanized antibody prepared in the Reference Examples 1 to 4 described later. (hereinafter this antibody is referred to as "humanized antibody")

[0118] In this Example, 10 kinds of preparations for injection were prepared as shown in Table 5.

Table 5

Preparation for injection	Buffer Solution	pH	humanized antibody
1	Acetic acid buffer solution 20mM	5	-
2	Acetic acid buffer solution 20mM	6	-
3	Acetic acid buffer solution 20mM	7	-
4	Acetic acid buffer solution 100mM	6	-
5	Citric acid buffer solution 20mM	5	-
6	Citric acid buffer solution 20mM	6	-
7	Citric acid buffer solution 20mM	7	-
8	Citric acid buffer solution 100mM	6	-
9	Acetic acid buffer solution 20mM	6	13.0mg/ml
10	Citric acid buffer solution 20mM	6	16.8mg/ml

[0119] These preparations for injection were quasi-aseptically prepared, kept in cold storage, and dispensed into a syringe for use at a time of administration.

Animal to be used and breeding environment

[0120] Rabbits were selected as the animal to be used, because their biological characteristics have been well studied, a large number of homogeneous rabbits are available, the size of a subcutaneous area around posterior auricular vein of a rabbit is suitable, as administration site, for administration and examination, and a macroscopic observation on a detriment site is easy. Male New Zealand white rabbits (Kbl:NZW) (12 week-old at purchase) were purchased from Kitayama Labes Co., Ltd., and 7 days acclimation breeding was conducted. During the acclimation breeding, the selection among them was made, considering general conditions and weights thereof. Then, 22 rabbits were selected for use. At a time of administration (13 week-old at administration), these animals had a weight of 2.6 to 3.3 kg.

[0121] These animals were each accommodated in a suspended aluminum cage having a three dimensions of 350 mm width × 500 mm depth × 350 mm height in an animal room with the conditions of temperature of  $24 \pm 2$  °C, humidity of  $55 \pm 10$  %, a light/dark cycle of 14 hours (from 5:00 to 19:00) of light per day, and 14 to 16 times ventilations per hour. Solid diet RC4 (available from Oriental Yeast Co., Ltd.) was served as feed through stainless steel feed supplier by an uninterrupted feeding method, and tap water was served as drinking water through an automatic water supply apparatus by an uninterrupted watering method. Both were served for animal's intake without restriction. For individual identification, the number of each animal was written on auricle (writing on not administration site but outside therefrom) by an oil marker pen, and for cage identification, a cage card was attached thereto.

[0122] During the examination period, in the analysis of breeding room environment, feed, and drinking water, there was recognized no defect which may cause any influence on the examination system.

## Setting of dose solution and group constitution

**[0123]** The dose volume of the injection preparations described in Table 5 was determined to be 0.2 mL/(administration site) in accordance with a local disorder examination which had already been conducted (Masao SUNAGA, Kazuhiro SHIMOMURA, Haruko KOIZUMI, Local Irritation Examination of Gadoteridol on rabbit at perivenous subcutaneous and intravenous administration, Preclin. Rep. Cent. Inst. Exp. Anim. 1992,18(1):47-57.). Each preparation for injection was administered to a group of 2 rabbits (4 auricle perivenous subcutaneous sites), and one of each group was autopsied 2 days after the administration, and the other was autopsied 4 days thereafter.

## Route of Administration

**[0124]** As an administration site of the preparation for injection, on a posterior auricular vein positioned almost at the center of auricle, a portion of the posterior auricular vein having less bifurcation of small veins was selected. For administration, a disposable syringe and needle (27G) were used. The needle was subcutaneously implanted toward a root of auricle along the vein, and then a slow single dose was provided in about 3 seconds. Further, in order to identify a needle implanting portion and a tip portion (injection portion) of the implanted needle, a marking was made on the side of the implanting portion and tip portion by an oil marker pen.

## Observation of administration site (macroscopic findings)

**[0125]** Observations were made on the administration site and the vicinity thereof once a day until 2 or 4 days had passed just after administration (counted from the date of next day of the administration). Then, the observed changes in size (long and short diameters) were measured by a vernier caliper (JIS Standard), obtaining the product of the long and short diameters as an area.

## Histopathological examination (histopathological findings)

**[0126]** After finishing the observation of the administration site during the period of 2 or 4 days just after the administration, body weight measurement was conducted for the calculation of anesthetic drug volume. The animals were euthanized under anesthesia of pentobarbital sodium (Nembutal available from Dainippon Pharmaceutical Co., Ltd) by exsanguinating from abdominal aorta. In passing, the anesthetic drug was carefully administered at posterior auricular vein of a root of auricle, and attention was paid not to cause influence on the evaluation of administration sites. Further, photos were taken on administration sites before the exsanguinations.

**[0127]** After collecting right and left auricles from a root of auricle to which the examination substance was administered, the collected auricles were fixed with neutral buffered formalin fixative. Thereafter, the examination substance-implanting portion of the right and left auricles, including a blood vessel adjacent to the administration site, was cut out in vertical direction to the run of the blood vessel, and paraffin embedded thin sectioning tissue preparations thereof were made by a conventional method. Hematoxylin and eosin (HE) stain was conducted thereon, and histopathological examination was conducted with a light

**[0128]** The irritancy according to histopathological examination was judged based on the following standard points.

(1) Bleeding was not considered as an index for irritancy but a change of administration technique.

(2) A slight change of tissue, such as slight cell density increase or lacuna formation of the administration sites, was not considered as an index for irritancy but a change of administration technique.

(3) In a case where there is other slight histological change, and vacuole formation of an epidermal cornification cell and epidermis thickening are observed, but is not considered as an index for irritancy. (These changes are considered to be expressed when there is a strong histological influence, and accordingly when other histological change is light, and vacuole formation of an epidermal cornification cell and epidermis thickening are observed this is not an index for irritancy.)

(4) When inflammatory cell infiltration and edema are more than slight, this is an index for irritancy.

(5) With respect to epidermis thickening, when inflammatory cell infiltration and edema are more than slight, this is an index for irritancy.

**[0129]** Then, biological evaluations were made on examination results, concerning difference among the groups or time-series changes. The evaluation results are shown in Table 6.



Table 6

Examined substances		2 days after administration				4 days after administration			
		right		left		right		left	
		macro- findings	histo- findings	irritancy	macro- findings	histo- findings	irritancy	macro- findings	histo- findings
Acetic acid buffer solution	20mM pH5	-	-	-	-	-	-	-	±
	20mM pH6	-	-	-	-	-	-	-	-
	20mM pH7	-	-	-	-	-	-	-	-
	100mM pH6	-	-	-	-	-	-	-	-
Citric acid buffer solution	20mM pH5	-	-	-	-	-	-	-	-
	20mM pH6	-	-	-	-	-	-	-	-
	20mM pH7	-	-	-	-	-	-	-	-
	100mM pH6	+	±	-	+	±	-	-	±
acetic acid buffer sol. containing 13.0 mg/mL of Humanized	20mM pH6	-	±	-	+	-	-	-	-
citric acid buffer sol. containing 16.8 mg/mL of Humanized	20mM pH6	+	+	+	+	+	+	++	+

Macroscopic findings - : not abnormal, ± : light erythema, + : erythema, ++ : erythema and swelling

Histopathological findings - : not abnormal, ± : bleeding, vacuole formation of an epidermal cornification cell, or slight change of tissue, ++ : inflammatory cell infiltration and edema

Irritancy - : negative, + : positive

**[0130]** As shown in Table 6, the solution of the humanized antibody (16.8 mg/Lm)/20 mmol/L of citric acid buffer solution (pH 6) provoked erythema and swelling by macroscopic observation, and light inflammatory cell infiltration and light edema by histopathological examination. Therefore, it was determined to have a local irritancy. In contrast to this, it was not recognized that the solution of the humanized antibody (13.0 mg/mL)/20 mmol/L of acetic acid buffer solution (pH 6) caused any macroscopic finding or histopathological finding, and thus it was determined to have no irritancy.

**[0131]** In view of the above results, it becomes clear that a pharmaceutical preparation for injection having a pain-easing action can be obtained by using acetic acid as buffer.

#### [REFERENCE EXAMPLE 1]

##### Preparation of hybridomas producing anti-PTHrP (1-34) mouse monoclonal antibody

**[0132]** Hybridomas capable of producing a monoclonal antibody against human PTHrP (1-34) (SEQ ID NO: 75), #23-57-154 and #23-57-137-1, were prepared as follows (see Sato, K. et al., J. Bone Miner. Res. 8, 849-860, 1993). The amino acid sequence of the human PTHrP (1-34) is shown in SEQ ID NO:75.

**[0133]** For use as an immunogen, PTHrP (1-34) (Peninsula) was conjugated with a carrier protein thyroglobulin using carbodiimide (Dojinn). The thyroglobulin-conjugated PTHrP (1-34) was dialyzed to obtain a solution having a protein concentration of 2 µg/ml. The resulting solution was mixed with Freund's adjuvant (Difco) at a mixing ratio of 1:1 to give an emulsion. This emulsion was injected to 16 female BALB/C mice 11 times subcutaneously at the back or intraperitoneally at a dose level of 100 µg/mouse for each injection, thereby immunizing the mice. For the priming immunization, Freund's complete adjuvant was used; while for the boosting immunization, Freund's incomplete adjuvant was used.

**[0134]** Each of the immunized mice was determined for its antibody titer in the serum in the following manner. That is, each of the mice was blood-drawn via its tail vein, and the anti-serum is separated from the blood. The anti-serum was diluted with a RIA buffer and mixed with <sup>125</sup>I-labeled PTHrP (1-34) to determine the binding activity. The mice that were confirmed to have a sufficiently increased titer were injected with PTHrP (1-34) without a carrier protein intraperitoneally at a dose level of 50 µg/mouse for the final immunization.

**[0135]** Three days after the final immunization, the mouse was sacrificed and the spleen was removed therefrom. The spleen cells were subjected to cell fusion with mouse myeloma cell line P3x63Ag8U.1 in accordance with a conventional known method using 50% polyethylene glycol 4000. The fused cells thus prepared were seeded to each well of eighty-five 96-well plates at a density of 2 x 10<sup>4</sup>/well. Hybridomas were screened in HAT medium as follows.

**[0136]** The screening of hybridomas was performed by determining the presence of PTHrP-recognition antibodies in the culture supernatant of the wells in which cell growth had been observed in HAT medium, by solid phase RIA method. The hybridomas were collected from the wells in which the binding ability to the PTHrP-recognition antibodies had been confirmed. The hybridomas thus obtained was suspended into RPMI-1640 medium containing 15% FCS supplemented with OPI-supplement (Sigma), followed by unification of the hybridomas by limiting dilution method. Thus, two types of hybridoma clones, #23-57-154 and #23-57-137-1, could be obtained, both which had a high binding ability to PTHrP (1-34).

**[0137]** Hybridoma clone #23-57-137-1 was designated "mouse-mouse hybridoma #23-57-137-1", and has been deposited under the terms of the Budapest Treaty on August 15, 1996 at the National Institute of Bioscience and Human-technology, Agency of Industrial Science and Technology, Japan (1-3, Higashi 1-chome, Tsukuba-shi, Ibaraki, Japan) under the accession No. FERM BP-5631.

#### [REFERENCE EXAMPLE 2]

##### Cloning of DNAs encoding V-regions of mouse monoclonal antibody against human PTHrP (1-34)

**[0138]** Cloning of DNAs encoding the V-regions of a mouse monoclonal antibody against human PTHrP (1-34), #23-57-137-1, was performed in the following manner.

##### (1) Preparation of mRNA

**[0139]** mRNA from hybridoma #23-57-137-1 was prepared using Quick Prep mRNA Purification Kit (Pharmacia Biotech). That is, cells of hybridoma #23-57-137-1 were fully homogenized with an extraction buffer, and mRNA was isolated and purified therefrom on an oligo(dT)-Cellulose Spun Column in accordance with the instructions included in the kit. The resulting solution was subjected to ethanol precipitation to obtain the mRNA as a precipitate. The mRNA precipitate was dissolved in an elution buffer.

## (2) Production and amplification of cDNA for gene encoding mouse H-chain V-region

## (i) Cloning of cDNA for #23-57-137-1 antibody H-chain V-region

5 **[0140]** A gene encoding H-chain V-region of the mouse monoclonal antibody against human PTHrP was cloned by 5'-RACE method (Frohman, M. A. et al., Proc. Natl. Acad. Sci. USA, 85, 8998-9002, 1988; Belyavsky, A. et al., Nucleic Acids Res. 17, 2919-2932, 1989). The 5'-RACE method was performed using 5'-Ampli FINDER RACE Kit (CLONETECH) in accordance with the instructions included in the kit. In this method, the primer used for synthesis of cDNA was MHC2 primer (SEQ ID NO: 1) which is capable of hybridizing to mouse H-chain C-region. The above-prepared mRNA (about 2 µg), which was a template for the cDNA synthesis, was mixed with MHC2 primer (10 pmoles).  
10 The resulting mixture was reacted with a reverse transcriptase at 52° C for 30 minutes to effect the reverse transcription of the mRNA into cDNA.

**[0141]** The resulting reaction solution was added with 6N NaOH to hydrolyze any RNA remaining therein (at 65° C for 30 min.) and then subjected to ethanol precipitation to isolate and purify the cDNA as a precipitate. The purified  
15 cDNA was ligated to Ampli FINDER Anchor (SEQ ID NO: 42) at the 5' end by reacting with T4 RNA ligase at 37° C for 6 hours and additionally at room temperature for 16 hours. As the primers for amplification of the cDNA by PCR method, Anchor primer (SEQ ID NO: 2) and MHC-G1 primer (SEQ ID NO: 3) (S.T. Jones, et al., Biotechnology, 9, 88, 1991) were used.

**[0142]** The PCR solution comprised (per 50 µl) 10 mM Tris-HCl (pH 8.3), 50 mM KCl, 0.25 mM dNTPs (dATP, dGTP, dCTP, dTTP), 1.5 mM MgCl<sub>2</sub>, 2.5 units of TaKaRa Taq (Takara Shuzo Co., Ltd.), 10 pmoles Anchor primer, and 1 µl of  
20 the reaction mixture of the cDNA to which MHC-G1 primer and Ampli FINDER Anchor primer had been ligated, over which mineral oil (50 µl) was layered. The PCR was performed on Thermal Cycler Model 480J (Perkin Elmer) for 30 cycles under the conditions: 94° C for 45 sec.; 60° C for 45 sec.; and 72° C for 2 min.

## 25 (ii) Cloning of cDNA for #23-57-137-1 antibody L-chain V-region

**[0143]** A gene encoding L-chain V-region of the mouse monoclonal antibody against human PTHrP was cloned by 5'-RACE method (Frohman, M. A. et al., Proc. Natl. Acad. Sci. USA, 85, 8998-9002, 1988; Belyavsky, A. et al., Nucleic Acids Res. 17, 2919-2932, 1989). The 5'-RACE method was performed using 5'-Ampli FINDER RACE Kit (CLONETECH)  
30 in accordance with the instructions included in the kit. In this method, oligo-dT primer was used as the primer for synthesizing cDNA. The above-prepared mRNA (about 2 µg), which was a template for the cDNA synthesis, was mixed with oligo-dT primer. The resulting mixture was reacted with a reverse transcriptase at 52° C for 30 min. to effect the reverse transcription of the mRNA into cDNA. The resulting reaction solution was added with 6N NaOH to hydrolyze any RNA remaining therein (at 65° C for 30 min.). The resulting solution was subjected to ethanol precipitation to isolate  
35 and purified the cDNA as a precipitate. The cDNA thus synthesized was ligated to Ampli FINDER Anchor at the 5' end by reacting with T4 RNA ligase at 37° C for 6 hours and additionally at room temperature for 16 hours.

**[0144]** A PCR primer MLC (SEQ ID NO: 4) was designed based on the conserved sequence of mouse L-chain λ chain C-region and then synthesized using 394 DNA/RNA Synthesizer (ABI). The PCR solution comprised (per 100 µl) 10 mM Tris-HCl (pH 8.3), 50 mM KCl, 0.25 mM dNTPs (dATP, dGTP, dCTP, dTTP), 1.5 mM MgCl<sub>2</sub>, 2.5 units of  
40 AmpliTaq (PERKIN ELMER), 50 pmoles of Anchor primer (SEQ ID NO: 2), and 1 µl of the reaction mixture of the cDNA to which MLC (SEQ ID NO: 4) and Ampli FINDER Anchor were ligated, over which mineral oil (50 µl) was layered. The PCR reaction was performed on Thermal Cycler Model 480J (Perkin Elmer) for 35 cycles under the conditions: 94° C for 45 sec.; 60° C for 45 sec.; and 72° C for 2 min.

## 45 (3) Purification and fragmentation of PCR products

**[0145]** Each of the DNA fragments amplified by PCR method described above was separated by agarose gel electrophoresis on a 3% Nu Sieve GTG agarose (FMC Bio. Products). For each of the H-chain V-region and the L-chain V-region, an agarose gel segment containing a DNA fragment of about 550 bp was excised from the gel. Each of the  
50 gel segments was subjected to purification of the DNA fragment of interest using GENECLEAN II Kit (BIO101) in accordance with the instructions included in the kit. The purified DNA was precipitated with ethanol, and the DNA precipitate was dissolved in 20 µl of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA. An aliquot (1 µl) of the DNA solution was digested with a restriction enzyme XmaI (New England Biolabs) at 37° C for 1 hour and further digested with a restriction enzyme EcoRI (Takara Shuzo Co., Ltd.) at 37° C for 1 hour. The digestion solution was  
55 extracted with phenol and chloroform and then precipitated with ethanol to collect the DNA.

**[0146]** In this manner, two DNA fragments containing a gene encoding mouse H-chain V-region and a gene encoding mouse L-chain V-region, respectively, were obtained, both which had an EcoRI recognition sequence on the 5' end and an XmaI recognition sequence on the 3' end.

**[0147]** The EcoRI-XmaI DNA fragments containing a gene encoding mouse H-chain V-region and a gene encoding mouse L-chain V-region, respectively, were separately ligated to pUC19 vector that had been digested with EcoRI and XmaI at 16° C for 1 hour using DNA Ligation Kit ver.2 (Takara Shuzo Co., Ltd.) in accordance with the instructions included in the kit. An aliquot (10 µl) of the ligation mixture was added to 100 µl of a solution containing competent cells of E. coli, JM 109 (Nippon Gene Co., Ltd.). The cell mixture was allowed to stand on ice for 15 min., at 42° C for 1 min. and additionally for 1 min. on ice. The resulting cell mixture was added with 300 µl of SOC medium (Molecular Cloning: A Laboratory Manual, Sambrook, et al., Cold Spring Harbor Laboratory Press, 1989) and then incubated at 37° C for 30 min. The resulting cell solution was plated on LB agar medium or 2xYT agar medium (Molecular Cloning: A Laboratory Manual, Sambrook, et al., Cold Spring Harbor Laboratory Press, 1989) containing either 100 or 50 µg/ml of ampicillin, 0.1 mM of IPTG and 20 µg/ml of X-gal, and then incubated at 37° C overnight. In this manner, E. coli transformants were prepared.

**[0148]** The transformants were cultured at 37° C overnight in 2 ml of LB or 2xYT medium containing either 100 or 50 µg/ml of ampicillin. The cell fraction was applied to Plasmid Extractor PI-100 (Kurabo Industries, Ltd.) or QIAprep Spin Plasmid Kit (QIAGEN) to give a plasmid DNA. The plasmid DNA was sequenced as follows.

#### (4) Sequencing of genes encoding mouse antibody V-regions

**[0149]** The nucleotide sequence of the cDNA coding region carried on the plasmid was determined in DNA Sequencer 373A (ABI; Perkin-Elmer) using Dye Terminator Cycle Sequencing Kit (Perkin-Elmer). M13 Primer M4 (Takara Shuzo Co., Ltd.) (SEQ ID NO: 5) and M13 Primer RV (Takara Shuzo Co., Ltd.) (SEQ ID NO: 6) were used as the primers for sequencing, and the nucleotide sequence was confirmed in the both directions.

**[0150]** The plasmid containing a gene encoding mouse H-chain V-region derived from hybridoma #23-57-137-1 was designated "MBC1H04", and the plasmid containing a gene encoding mouse L-chain V-region derived from hybridoma #23-57-137-1 was designated "MBC1L24". The nucleotide sequences (including the corresponding amino acids sequences) of the gene encoding the mouse #23-57-137-1 antibody-derived H-chain V-region in plasmid MBC1H04 and the gene encoding the mouse #23-57-137-1 antibody-derived L-chain V-region in plasmid MBC1L24 were shown in SEQ. ID Nos: 57 and 65, respectively. The amino acid sequences of the polypeptides for the H-chain V-region and the L-chain V-region were shown in SEQ. ID NOs: 46 and 45, respectively.

**[0151]** The E. coli strain containing plasmid MBC1H04 and the E. coli strain containing plasmid MBC1L24 were designated "Escherichia coli JM109 (MBC1H04)" and "Escherichia coli JM 109 (MBC1L24)", respectively. These E. coli strains have been deposited under the terms of the Budapest Treaty at the National Institute of Bioscience and Human-Technology, Agency of Industrial Science and Technology, Japan (1-3, Higashi 1-chome, Tsukuba-shi, Ibaraki, Japan) on August 15, 1996, under the Accession No. FERM BP-5628 for Escherichia coli JM109 (MBC1H04) and FERM BP-5627 for Escherichia coli JM109 (MBC1L24), respectively.

#### (5) Determination of CDRs of mouse monoclonal antibody

#23-57-137-1 against human PTHrP

**[0152]** The H-chain V-region and the L-chain V-region have general structures similar to each other, each of which has four framework regions (FRs) linked through three hypervariable regions (i.e., complementarity determining regions; CDRs). The amino acid sequences of the FRs are relatively well conserved, while the amino acid sequence of the CDRs have an extremely high variability (Kabat, E.A. et al., "Sequence of Proteins of Immunological Interest", US Dept. Health and Human Services, 1983).

**[0153]** In view of these facts, the homology in amino acid between the V-regions of the mouse monoclonal antibody against human PTHrP was determined with reference to the database of amino acid sequences of antibodies established by Kabat et al. Thus, the CDRs of the V-regions were determined as shown in Table 7.

**[0154]** The amino acid sequences of CDRs 1-3 in the L-chain V-region are shown in SEQ ID Nos: 59 to 61, respectively; and the amino acid sequences of CDRs 1-3 in the H-chain V-region are shown in SEQ ID Nos: 62 to 64, respectively.

Table 7

V-region	SEQ ID NO.	CDR1	CDR2	CDR3
H-chain V-region	57	31-35	50-66	99-107
L-chain V-region	65	23-34	50-60	93-105

## [REFERENCE EXAMPLE 3] Construction of Chimeric Antibody

## (1) Construction of chimeric antibody H-chain

## (i) Construction of H-chain V-region

**[0155]** To ligate to an expression vector carrying a genomic DNA of human H-chain C-region C  $\gamma$  1, the cloned DNA encoding mouse H-chain V-region was modified by PCR method. A backward primer MBC1-S1 (SEQ ID NO: 7) was designed to hybridize to a DNA sequence encoding the 5' region of the leader sequence of the V-region and to have both a Kozak consensus sequence (Kozak, M. et al., J. Mol. Biol., 196, 947-950, 1987) and a HindIII-recognition sequence. A forward primer MBC1-a (SEQ ID NO: 8) was designed to hybridize to a DNA sequence encoding the 3' region of the J region and to have both a donor splice sequence and a BamHI-recognition sequence. The PCR reaction was performed using TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) and a buffer appended thereto. The PCR solution comprised (per 50  $\mu$ l) 0.07  $\mu$ g of plasmid MBC1H04 as a template DNA, 50 pmoles of MBC1-a and 50 pmoles of MBC1-S1 as primers, 2.5U of TaKaRa Ex Taq and 0.25 mM dNTPs in the buffer, over which 50  $\mu$ l of mineral oil was layered. The PCR was run for 30 cycles under the conditions: 94° C for 1 min.; 55° C for 1 min.; 72° C for 2 min. The DNA fragments thus amplified by the PCR method were separated by agarose gel electrophoresis on a 3% Nu Sieve GTG Agarose (FMC Bio. Products).

**[0156]** Then, an agarose gel segment containing a DNA fragment of 437 bp was excised, and the DNA fragment was purified therefrom using GENECLAN II Kit (BIO101) in accordance with the instructions included in the kit. The purified DNA was collected by ethanol precipitation, and then dissolved in 20  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA. An aliquot (1  $\mu$ l) of the resulting DNA solution was digested with restriction enzymes BamHI and HindIII (Takara Shuzo Co., Ltd.) at 37° C for 1 hour. The digestion solution was extracted with phenol and chloroform and then precipitated with ethanol to collect the DNA of interest.

**[0157]** The obtained HindIII-BamHI DNA fragment, which containing a gene encoding the mouse H-chain V-region, was subcloned into pUC19 vector that had been digested with HindIII and BamHI. The resulting plasmid was sequenced on DNA Sequencer 373A (Perkin-Elmer) using M13 Primer M4 and M13 Primer RV as primers and Dye Terminator Cycle Sequencing Kit (Perkin-Elmer). As a result, a plasmid which carried a gene of correct nucleotide sequence encoding the mouse H-chain V-region derived from hybridoma #23-57-137-1 and had a HindIII-recognition sequence and a Kozak sequence on its 5' region and a BamHI-recognition sequence on its 3' region was obtained, which was designated "MBC1H/pUC19".

## (ii) Construction of H-chain V-region for preparation of cDNA-type of mouse-human chimeric H-chain

**[0158]** To ligate to cDNA of the human H-chain C-region C  $\gamma$  1, the DNA encoding the mouse H-chain V-region constructed as described above was modified by PCR method. A backward primer MBC1HVS2 (SEQ ID NO: 9) for the V-region was designed to cause the replacement of the second amino acid (asparagine) of the sequence encoding the front part of the leader sequence of the H-chain V-region by glycine and to have a Kozak consensus sequence (Kozak, M. et al., J. Mol. Biol., 196, 947-950, 1987) and HindIII- and EcoRI-recognition sequences. A forward primer MBC1HVR2 (SEQ ID NO: 10) for the H-chain V-region was designed to hybridize to a DNA sequence encoding the 3' region of the J region, to encoding the 5' region of the C-region and to have ApaI- and SmaI-recognition sequences.

**[0159]** The PCR reaction was performed using TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) and a buffer appended thereto. The PCR solution comprised (per 50  $\mu$ l) 0.6  $\mu$ g of plasmid MBC1H/pUC19 as a template DNA, 50 pmoles of MBC1HVS2 and 50 pmoles of MBC1HVR2 as primers, 2.5U of TaKaRa Ex Taq and 0.25 mM of dNTPs in the buffer, over which 50  $\mu$ l of mineral oil was layered. The PCR reaction was run for 30 cycles under the conditions: 94° C for 1 min.; 55° C for 1 min.; 72° C for 1 min. The DNA fragments amplified by the PCR reaction were separated by agarose gel electrophoresis on a 1 % Sea Kem GTG Agarose (FMC Bio. Products). Then, an agarose gel segment containing a DNA fragment of 456 bp was excised and the DNA fragment was purified therefrom using GENECLAN II Kit (BIO101) in accordance with the instructions included in the kit. The purified DNA was precipitated with ethanol and then dissolved in 20  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

**[0160]** The resulting DNA solution (1  $\mu$ g) was digested with restriction enzymes EcoRI and SmaI (Takara Shuzo Co., Ltd.) at 37° C for 1 hour. The digestion solution was extracted with phenol and chloroform and then precipitated with ethanol to collect the DNA. The obtained EcoRI-SmaI DNA fragment, which containing a gene encoding the mouse H-chain V-region, was subcloned into pUC19 vector that had been digested with EcoRI and SmaI. The resulting plasmid was sequenced on DNA Sequencer 373A (Perkin-Elmer) using M13 Primer M4 and M13 Primer RV, and Dye Terminator Cycle Sequencing Kit (Perkin-Elmer). As a result, a plasmid which contained a gene of correct nucleotide sequence encoding mouse H-chain V-region derived from hybridoma #23-57-137-1 and had EcoRI- and HindIII-recognition sequences and a Kozak sequence on its 5' region and ApaI- and SmaI-recognition sequences on its 3' region was obtained,

which was designated "MBC1Hv/pUC19".

(iii) Construction of expression vector for chimeric antibody H-chain

5 **[0161]** cDNA containing the DNA for human antibody H-chain C-region C $\gamma$ 1 was prepared as follows. mRNA was prepared from a CHO cell into which both an expression vector DHFR- $\Delta$ E-RVh-PM-1-f (see WO 92/19759) encoding the genomic DNAs of humanized PM1 antibody H-chain V-region and human antibody H-chain C-region IgG1 (N. Takahashi et al., Cell 29, 671-679, 1982) and an expression vector RV1-PM1a (see WO 92/19759) encoding the genomic DNAs of humanized PM1 antibody L-chain V-region and human antibody L-chain  $\kappa$  chain C-region had been introduced. Using the mRNA, cDNA containing the humanized PM1 antibody H-chain V-region and the human antibody C-region C $\gamma$ 1 was cloned by RT-PCR method, and then subcloned into plasmid pUC19 at the HindIII-BamHI site. After sequencing, a plasmid which had the correct nucleotide sequence was obtained, which was designated "pRVh-PM1f-cDNA".

15 **[0162]** An expression vector DHFR- $\Delta$ E-RVh-PM-1-f in which both a HindIII site located between SV40 promoter and a DHFR gene and an EcoRI site located between EF-1  $\alpha$  promoter and a humanized PM1 antibody H-chain V-region gene had been deleted, was prepared for the construction of an expression vector for cDNA containing the humanized PM1 antibody H-chain V-region gene and the human antibody C-region C $\gamma$ 1 gene.

20 **[0163]** The plasmid obtained (pRVh-PM1f-cDNA) was digested with BamHI, blunt-ended with Klenow fragment, and further digested with HindIII, thereby obtaining a blunt-ended HindIII-BamHI fragment. The blunt-ended HindIII-BamHI fragment was ligated to the above-mentioned HindIII site- and EcoRI site-deleted expression vector DHFR- $\Delta$ E-RVh-PM1-f that had been digested with HindIII and BamHI. Thus, an expression vector RVh-PM1f-cDNA was constructed which contained cDNA encoding the humanized PM1 antibody H-chain V-region and the human antibody C-region C $\gamma$ 1.

25 **[0164]** The expression vector RVh-PM1f-cDNA containing the cDNA encoding the humanized PM1 antibody H-chain V-region and the human antibody C-region C $\gamma$ 1 was digested with Apal and BamHI, and a DNA fragment containing the H-chain C-region was collected therefrom. The resulting DNA fragment was introduced into the plasmid MBC1Hv/pUC 19 that had been digested with Apal and BamHI. The plasmid thus prepared was designated "MBC1HcDNA/pUC19". This plasmid contained cDNA encoding the mouse antibody H-chain V-region and the human antibody C-region C $\gamma$ 1, and had EcoRI- and HindIII-recognition sequences on its 5' region and a BamHI-recognition sequence on its 3' region.

30 **[0165]** The plasmid MBC1HcDNA/pUC19 was digested with EcoRI and BamHI to give a DNA fragment comprising a nucleotide sequence encoding the chimeric antibody H-chain. The resulting DNA fragment was introduced into an expression vector pCOS1 that had been digested with EcoRI and BamHI, thereby giving an expression vector for the chimeric antibody, which was designated "MBC1HcDNA/pCOS1". Here, the expression vector pCOS1 was constructed using HEF-PMh-gy1 (see WO 92/19759) by deleting therefrom an antibody gene by digestion with EcoRI and SmaI, and then ligating it to EcoRI-NotI-BamHI Adaptor (Takara Shuzo Co., Ltd.).

35 **[0166]** For preparing a plasmid for the expression in a CHO cell, the plasmid MBC1HcDNA/pUC19 was digested with EcoRI and BamHI to obtain a DNA fragment containing a gene for the chimeric antibody H-chain. The DNA fragment was then introduced into an expression plasmid pCHO1 that had been digested with EcoRI and BamHI to give an expression plasmid for the chimeric antibody, which was designated "MBC1HcDNA/pCHO1". Here, the expression vector pCHO1 was constructed using DHFR- $\Delta$ E-rvH-PM1-f (see WO 92/19759) by deleting therefrom an antibody gene by digestion with EcoRI and SmaI, and then ligating it to EcoRI-NotI-BamHI Adaptor (Takara Shuzo Co., Ltd.).

(2) Construction of human L-chain C-region

45 (i) Preparation of cloning vector

50 **[0167]** To construct pUC19 vector containing a gene for human L-chain C-region, a HindIII site-deleted pUC19 vector was prepared. pUC19 vector (2  $\mu$ g) was digested in 20  $\mu$ l of a reaction solution containing 20 mM Tris-HCl (pH 8.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 100 mM KCl, 8 U of HindIII (Takara Shuzo Co., Ltd.) at 37° C for 1 hour. The resulting digestion solution was extracted with phenol and chloroform, and then subjected to ethanol precipitation to collect the DNA of interest.

55 **[0168]** The DNA collected was reacted in 50  $\mu$ l of a reaction solution containing 50 mM Tris-HCl (pH 7.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 100 mM NaCl, 0.5 mM dNTPs and 6U of Klenow fragment (GIBCO BRL) at room temperature for 20 min., thereby rendering the terminal ends of the DNA blunt. This reaction mixture was extracted with phenol and chloroform and then subjected to ethanol precipitation to collect the vector DNA.

**[0169]** The vector DNA thus collected was reacted in 10  $\mu$ l of a reaction solution containing 50 mM Tris-HCl (pH 7.6), 10 mM MgCl<sub>2</sub>, 1 mM ATP, 1 mM DTT, 5% (v/v) polyethylene glycol-8000 and 0.5 U of T4 DNA ligase (GIBCO BRL) at

16 °C for 2 hours, to cause self-ligation of the vector DNA. The reaction solution (5 µl) was added to 100 µl of a solution containing competent cells of *E. coli*, JM109 (Nippon Gene Co., Ltd.), and the resulting solution was allowed to stand on ice for 30 min., at 42 °C for 1 min., and additionally on ice for 1 min. SOC culture medium (500 µl) was added to the reaction solution and then incubated at 37 °C for 1 hour. The resulting solution was plated on 2 × YT agar medium (containing 50 µg/ml of ampicillin) on which X-gal and IPTG had been applied (Molecular Cloning: A Laboratory Manual, Sambrook, et al., Cold Spring Harbor Laboratory Press, 1989), and then cultured at 37 °C overnight, thereby obtaining a transformant.

[0170] The transformant was cultured in 2 × YT medium (20 ml) containing ampicillin (50 µg/ml) at 37 °C overnight. From the cell fraction of the culture medium, a plasmid DNA was isolated and purified using Plasmid Mini Kit (QIAGEN) in accordance with the instructions included in the kit. The purified plasmid was digested with HindIII. The plasmid that was confirmed to have a HindIII site-deletion was designated "pUC19 ΔHindIII".

## (ii) Construction of DNA encoding human L-chain λ chain C-region

[0171] Human antibody L-chain λ chain C-region is known to have at least four isotypes including Mcg<sup>+</sup>Ke<sup>+</sup>Oz<sup>-</sup>, Mcg<sup>-</sup>Ke<sup>-</sup>Oz<sup>-</sup>, Mcg<sup>-</sup>Ke<sup>+</sup>Oz<sup>+</sup> and Mcg<sup>+</sup>Ke<sup>+</sup>Oz<sup>+</sup> (P. Dariavach, et al., Proc. Natl. Acad. Sci. USA, 84, 9074-9078, 1987). A search was made for a human antibody L-chain λ chain C-region homologous to the #23-57-137-1 mouse L-chain λ chain C-region from the EMBL database. As a result, it was found that the isotype Mcg<sup>+</sup>Ke<sup>+</sup>Oz<sup>-</sup> of the human antibody L-chain λ chain (Accession No. X57819) (P. Dariavach, et al., Proc. Natl. Acad. Sci. USA, 84, 9074-9078, 1987) showed the highest degree of homology to the #23-57-137-1 mouse L-chain λ chain C-region, with a 64.4% homology in terms of amino acid sequence and a 73.4% homology in terms of nucleotide sequence.

[0172] Then, a gene encoding the human antibody L-chain λ chain C-region was constructed by PCR method. The primers for the PCR were synthesized using 394 DNA/RNA Synthesizer (ABI). The synthesized primers were as follows: HLAMB1 (SEQ ID NO: 11) and HLAMB3 (SEQ ID NO: 13), both having a sense DNA sequence; and HLAMB2 (SEQ ID NO: 12) and HLAMB4 (SEQ ID NO: 14), both having an antisense DNA sequence; each primer containing a complementary sequence of 20-23 bp on the both terminal ends.

[0173] External primers HLAMBS (SEQ ID NO: 15) and HLAMBR (SEQ ID NO: 16) had sequences homologous to the primers HLAMB1 and HLAMB4, respectively. HLAMBS contained EcoRI-, HindIII- and B1nI-recognition sequences, and HLAMBR contained an EcoRI-recognition sequence. In the first-round PCR reaction, the reactions between HLAMB1 and HLAMB2 and between HLAMB3 and HLAMB4 were performed. After the reactions were completed, both of the resulting PCR products were mixed in equivalent quantities, and then assembled in the second-round PCR reaction. The reaction solution was added with the external primers HLAMBS and HLAMBR. This reaction mixture was subjected to the third-round PCR reaction to amplify the full length DNA.

[0174] Each PCR reaction was performed using TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) in accordance with the instructions included in the kit. In the first-round PCR reaction, 100 µl of either a reaction solution containing 5 pmoles of HLAMB1, 0.5 pmole of HLAMB2 and 5U of TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) or a reaction solution containing 0.5 pmole of HLAMB3, 5 pmoles of HLAMB4 and 5U of TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) was used, over which 50 µl of mineral oil was layered. The PCR reaction was run for 5 cycles under the conditions: 94° C for 1 min., 60° C for 1 min. and 72° C for 1 min.

[0175] In the second-round PCR reaction, a mixture of both the reaction solutions (50 µl each) was used, over which 50 µl of mineral oil was layered. The PCR reaction was run for 3 cycles under the conditions: 94° C for 1 min., 60° C for 1 min. and 72° C for 1 min.

[0176] In the third-round PCR reaction, the reaction solution to which the external primers HLAMBS and HLAMBR (50 pmoles each) were added was used. The PCR reaction was run for 30 cycles under the conditions: 94° C for 1 min., 60° C for 1 min. and 72° C for 1 min.

[0177] The DNA fragment obtained by the third-round PCR reaction was subjected to electrophoresis on a 3% low-melting agarose gel (NuSieve GTG Agarose, FMC), and separated and purified from the gel using GENECLAN II Kit (BIO101) in accordance with the instructions included in the kit.

[0178] The DNA fragment obtained was digested in a reaction solution (20 µl) containing 50 mM Tris-HCl (pH 7.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 100 mM NaCl and 8U of EcoRI (Takara Shuzo Co., Ltd.) at 37° C for 1 hour. The digestion solution was extracted with phenol and chloroform, and the DNA was collected therefrom by the ethanol precipitation. The DNA was dissolved in a solution (8 µl) containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

[0179] The above-prepared plasmid pUC19 ΔHindIII (0.8 µg) was digested with EcoRI in the same manner as set forth above. The digestion solution was subjected to phenol/chloroform extraction and then ethanol precipitation, thereby giving a digested plasmid pUC19 ΔHindIII. The digested plasmid was reacted in a reaction solution (50 µl) containing 50 mM Tris-HCl (pH 9.0), 1 mM MgCl<sub>2</sub> and alkaline phosphatase (*E. coli* C75; Takara Shuzo Co., Ltd.) at 37° C for 30 min. to dephosphorylate (i.e., BAP-treat) the plasmid. The reaction solution was subjected to phenol/chloroform extraction, and the DNA was collected therefrom by ethanol precipitation. The DNA thus obtained was dissolved in a

solution (10 µl) containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

[0180] The BAP-treated plasmid pUC19 ΔHindIII (1 µl) was ligated to the above-obtained PCR product (4 µl) using DNA Ligation Kit Ver.2 (Takara Shuzo Co., Ltd.). The resulting plasmid was introduced into a competent cell of *E. coli*, JM109, to give a transformant. The transformant was cultured overnight in 2xYT medium (2 ml) containing 50 µg/ml

of ampicillin. From the cell fraction, the plasmid was isolated using QIAprep Spin Plasmid Kit (QIAGEN).

[0181] The plasmid obtained was sequenced for the cloned DNA part. The sequencing was performed on 373A DNA Sequencer (ABI) using M13 Primer M4 and M13 Primer RV (Takara Shuzo Co., Ltd.). As a result, it was found that the cloned DNA had a 12-bp deletion therein. The plasmid was designated "C λ ΔpUC19". Then, for making up for the deleted part, primers HCLMS (SEQ ID NO: 17) and HCLMR (SEQ ID NO: 18) were newly synthesized, and a DNA of

correct sequence was reconstructed using these primers by PCR method.

[0182] In the first-round PCR reaction, the plasmid C λ ΔpUC19 having the DNA deletion therein was used as a template, and the reaction was performed with each of the primer sets of HLAMBS and HCLMS and HCLMS and HLAMB4. The PCR products were purified separately. In the second-round PCR reaction, the PCR products were assembled together. In the third-round PCR reaction, the reaction product of the second-round PCR reaction was

added with external primers HLAMBS and HLAMB4 and amplified to give the full length DNA.

[0183] In the first-round PCR reaction, a reaction solution (100 µl) containing 0.1 µg of C λ ΔpUC19 as a template, either 50 pmoles of each of the primers HLAMBS and HCLMR or 50 pmoles of each of the primers HCLMS and HLAMB4, and 5U of TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) was used, over which 50 µl of mineral oil was layered. The PCR reaction was run for 30 cycles under the conditions: 94° C for 1 min., 60° C for 1 min. and 72° C for 1 min.

[0184] The PCR products of the first-round PCR reaction, HLAMBS-HCLMR (236 bp) and HCLMS-HLAMB4 (147 bp), were subjected to electrophoresis separately on a 3% low-melting agarose gel to isolate the DNA fragments. The DNA fragments were collected and purified from the gels using GENE CLEAN II Kit (BIO101). In the second-round PCR reaction, 20 µl of a reaction solution containing 40 ng of each of the purified DNA fragments and 1U of TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) was used, over which 25 µl of mineral oil was layered. The PCR reaction was run for

5 cycles under the conditions: 94° C for 1 min., 60° C for 1 min. and 72° C for 1 min.

[0185] In the third-round PCR reaction, 100 µl of a reaction solution containing 2 µl of the reaction solution obtained by the second-round PCR reaction, 50 pmoles of each of external primers HLAMBS and HLAMB4 and 5U of TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) was used, over which 50 µl of mineral oil was layered. The PCR reaction was run for 30 cycles under the conditions: 94° C for 1 min., 60° C for 1 min. and 72° C for 1 min., thereby obtaining a DNA fragment of 357 bp (the third PCR product). The DNA fragment was subjected to electrophoresis on a 3% low-melting agarose gel to isolate the DNA fragment. The resulting DNA fragment was collected and purified using GENE CLEAN II Kit (BIO101).

[0186] An aliquot (0.1 µg) of the DNA fragment thus obtained was digested with EcoRI, and then subcloned into plasmid pUC19 ΔHindIII that had been BAP-treated. The resulting plasmid was introduced into a competent cell of *E. coli*, JM109, to form a transformant. The transformant was cultured overnight in 2 ml of 2xYT medium containing 50 µg/ml of ampicillin. From the cell fraction, the plasmid was isolated and purified using QIAprep Spin Plasmid Kit (QIAGEN).

[0187] The purified plasmid was sequenced on 373A DNA Sequencer (ABI) using M13 Primer M4 and M13 Primer RV (Takara Shuzo Co., Ltd.). The plasmid that was confirmed to have the correct nucleotide sequence without any deletion was designated "C λ /pUC19".

### (iii) Construction of gene encoding human L-chain κ chain C-region

[0188] A DNA fragment encoding the L-chain κ, chain C-region was cloned from plasmid HEF-PM1k-gk (WO 92/19759) by PCR method. A forward primer HKAPS (SEQ ID NO: 19) was designed to contain EcoRI-, HindIII and BlnI-recognition sequences, and a backward primer HKAPA (SEQ ID NO: 20) was designed to contain an EcoRI-recognition sequence. These primers were synthesized on 394 DNA/RNA Synthesizer (ABI).

[0189] A PCR reaction was performed using 100 µl of a reaction solution containing 0.1 µg of plasmid HEF-PM1k-gk as a template, 50 pmoles of each of primers HKAPS and HKAPA and 5U of TaKaRa Ex Taq (Takara Shuzo Co., Ltd.), over which 50 µl of mineral oil was layered. The PCR reaction was run for 30 cycles under the conditions: 94° C for 1 min., 60° C for 1 min. and 72° C for 1 min., thereby giving a PCR product of 360 bp. The DNA fragment was isolated and purified by electrophoresis on a 3% low-melting agarose, and then collected and purified using GENE CLEAN II Kit (BIO101).

[0190] The DNA fragment thus obtained was digested with EcoRI, and then cloned into plasmid pUC19 (HindIII that had been BAP-treated). The resulting plasmid was introduced into a competent cell of *E. coli*, JM109, to form a transformant. The transformant was cultured overnight in 2 ml of 2xYT medium containing 50 µg/ml of ampicillin. From the cell fraction, the plasmid was purified using QIAprep Spin Plasmid Kit (QIAGEN).

[0191] The purified plasmid was sequenced on 373A DNA Sequencer (ABI) using M13 Primer M4 and M13 Primer



RV (Takara Shuzo Co., Ltd.). The plasmid that was confirmed to have the correct nucleotide sequence was designated "C  $\kappa$ /pUC19".

### (3) Construction of chimeric antibody L-chain expression vector

**[0192]** An expression vector for the chimeric #23-57-137-1 antibody L-chain was constructed. A gene encoding #23-57-137-1L-chain V-region was ligated to the HindIII-B1nI site (located just in front of the human antibody C-region) of each of the plasmids C  $\lambda$  /pUC19 and C  $\kappa$  /pUC19, thereby obtaining pUC19 vectors that contained the DNAs encoding the chimeric #23-57-137-1 antibody L-chain V-region and either of the L-chain  $\lambda$  chain C-region or the L-chain  $\kappa$  region C-region, respectively. Each of the resulting vectors was then digested with EcoRI to separate the gene for the chimeric antibody L-chain. The gene was subcloned into HEF expression vector.

**[0193]** That is, a DNA fragment encoding #23-57-137-1 antibody L-chain V-region was cloned from plasmid MBC1L24 by PCR method. Primers used in the PCR method were separately synthesized using 394 DNA/RNA Synthesizer (ABI). A backward primer MBCCHL1 (SEQ ID NO: 21) was designed to contain a HindIII-recognition sequence and a Kozak sequence (Kozak, M. et al., J. Mol. Biol. 196, 947-950, 1987), and a forward primer MBCCHL3 (SEQ ID NO: 22) was designed to contain BglII- and RcoRI-recognition sequences.

**[0194]** The PCR reaction was performed using 100  $\mu$ l of a reaction solution containing 10 mM Tris-HCl (pH 8.3), 50 mM KCl, 1.5 mM MgCl<sub>2</sub>, 0.2 mM dNTPs, 0.1  $\mu$ g MBC1 L24, 50 pmols of each of primers MBCCHL1 and MBCCHL3 and 1  $\mu$ l of AmpliTaq (PERKIN ELMER), over which 50  $\mu$ l of mineral oil was layered. The PCR reaction was run for 30 cycles under the conditions: 94° C for 45 sec., 60° C for 45 sec. and 72° C for 2 min.

**[0195]** A PCR product of 444 bp was electrophoresed on a 3% low-melting agarose gel, and collected and purified using GENECLAN II Kit (BIO101). The purified PCR product was dissolved in 20  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA. The PCR product (1  $\mu$ l) was digested in 20  $\mu$ l of a reaction solution containing 10 mM Tris-HCl (pH 7.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 50 mM NaCl, 8U of HindIII (Takara Shuzo Co., Ltd.) and 8U of EcoRI (Takara Shuzo Co., Ltd.) at 37° C for 1 hour. The digestion solution was subjected to phenol/chloroform extraction, and the DNA of interest was collected therefrom by ethanol precipitation. The DNA was dissolved in 8  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

**[0196]** In the same manner, plasmid pUC19 (1  $\mu$ g) was digested with HindIII and EcoRI, and subjected to phenol/chloroform extraction and then ethanol precipitation. The obtained digested plasmid was BAP-treated with alkaline phosphatase (E. coli C75; Takara Shuzo Co., Ltd.). The resulting reaction solution was extracted with phenol and chloroform, and the DNA was collected therefrom by ethanol precipitation. The DNA was dissolved in 10  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

**[0197]** The BAP-treated plasmid pUC19 (1  $\mu$ l) was ligated to the above-obtained PCR product (4  $\mu$ l) using DNA Ligation Kit Ver. 2 (Takara Shuzo Co., Ltd.). The resulting plasmid was introduced into a competent cell of E. coli, JM109 (Nippon Gene Co., Ltd.), in the same manner as set forth above, to form a transformant. The transformant was plated on 2xYT agar medium containing 50  $\mu$ g/ml of ampicillin and cultured at 37° C overnight. The resulting transformant was then cultured at 37° C overnight in 2 ml of 2xYT medium containing 50  $\mu$ g/ml of ampicillin. From the cell fraction, the plasmid was purified using QIAprep Spin Plasmid Kit (QIAGEN). After determining the nucleotide sequence, the plasmid that was confirmed to have the correct nucleotide sequence was designated "CHL/pUC19".

**[0198]** Each of plasmids C  $\lambda$  /pUC19 and C  $\kappa$  /pUC19 (1  $\mu$ g each) was digested in 20  $\mu$ l of a reaction solution containing 20 mM Tris-HCl (pH 8.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 100 mM KCl, 8U of HindIII (Takara Shuzo Co., Ltd.) and 2U of B1nI (Takara Shuzo Co., Ltd.) at 37° C for 1 hour. The digestion solution was extracted with phenol and chloroform, and the DNA was collected therefrom by ethanol precipitation. The DNA was BAP-treated at 37° C for 30 min. The reaction solution was extracted with phenol and chloroform, and the DNA was collected therefrom by ethanol precipitation. The DNA was dissolved in 10  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

**[0199]** The plasmid CHL/pUC19 (8  $\mu$ g) that contained DNA encoding #23-57-137-1L-chain V-region was digested with HindIII and B1nI in the same manner as set forth above to give a DNA fragment of 409 bp. The DNA fragment was electrophoresed on a 3% low-melting agarose gel, and then collected and purified from the gel using GENECLAN II Kit (BIO101). The DNA was dissolved in 10  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

**[0200]** The DNA for L-chain V-region DNA (4  $\mu$ l) was subcloned into 1  $\mu$ l of each of the BAP-treated plasmids C  $\lambda$  /pUC19 and C  $\kappa$  /pUC19, and then introduced into a competent cell of E. coli, JM109, to form a transformant. The transformant was cultured overnight in 3 ml of 2xYT medium containing 50  $\mu$ g/ml of ampicillin. From the cell fraction, the plasmid was isolated and purified using QIAprep Spin Plasmid Kit (QIAGEN). The two plasmids thus prepared were designated "MBC1L( $\lambda$ )/pUC19" and "MBC1L( $\kappa$ )/pUC19", respectively.

**[0201]** Each of plasmids MBC1L( $\lambda$ )/pUC19 and MBC1L( $\kappa$ )/pUC19 was digested with EcoRI and then subjected to electrophoresis on a 3% low-melting agarose gel. A DNA fragment of 743 bp was isolated and purified from the gel using GENECLANII Kit (BIO101), and then dissolved in 10  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

**[0202]** An expression vector (plasmid HEF-PM1k-gk) (2.7 µg) was digested with EcoRI and then extracted with phenol and chloroform, and the DNA was collected therefrom by ethanol precipitation. The DNA fragment was BAP-treated, and then subjected to electrophoresis on a 1 % low-melting agarose gel. From the gel, a DNA fragment of 6561 bp was isolated and purified therefrom using GENECLANII Kit (BIO101). The purified DNA fragment was dissolved in

10 µl of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

**[0203]** BAP-treated HEF vector (2 µl) was ligated to an EcoRI fragment (3 µl) of each of plasmid MBC1L(λ)/pUC19 and MBC1L(κ)/pUC19. The ligation product was introduced into a competent cell of *E. coli* JM109, to form a transformant. The transformant was cultured in 2 ml of 2xYT medium containing 50 µg/ml of ampicillin. From the cell fraction, the plasmid was purified using QIAprep Spin Plasmid Kit (QIAGEN).

**[0204]** The purified plasmid was digested in 20 µl of a reaction solution containing 20 mM Tris-HCl (pH 8.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 100 mM KCl, 8U of HindIII (Takara Shuzo Co., Ltd.) and 2 U of PvuI (Takara Shuzo Co., Ltd.) at 37° C for 1 hour. This reaction gave digestion fragments of 5104/2195 bp if the fragment was inserted in the correct orientation, or gave digestion fragments of 4378/2926 bp if the fragment was inserted in the reverse orientation. The plasmid that was confirmed to have the fragment in the correct orientation was designated "MBC1L(λ)/neo" for plasmid MBC1L(λ)/pUC19 or "MBC1L(κ)/neo" for plasmid MBC1L(κ)/pUC19.

#### (4) Transfection of COS-7 cell

**[0205]** To evaluate the antigen-binding activity and the neutralizing activity of the chimeric antibodies, the expression plasmids prepared above were separately expressed transiently in a COS-7 cell.

**[0206]** The transient expression of the chimeric antibodies was performed using each of the combinations of plasmids MBC1HcDNA/pCOS1 and MBC1L(λ)/neo and plasmids MBC1HcDNA/pCOS1 and MBC1L(κ)/neo, by co-transfecting a COS-7 cell with the plasmids by electroporation using Gene Pulser (Bio Rad). That is, the plasmids (10 µg each) were added to a COS-7 cell suspension (0.8 ml; 1 x 10<sup>7</sup> cells/ml) in PBS(-). The resulting solution was applied with pulses at an electrostatic capacity of 1,500V and 2 µF to cause electroporation. After 10 min. of recovery period at room temperature, the electroporated cells were suspended in DMEM medium (GIBCO) containing 2% Ultra Low IgG fetal calf serum (GIBCO), and then cultured using a 10-cm culture dish in a CO<sub>2</sub> incubator. After culturing for 72 hours, a culture supernatant was collected and centrifuged to remove cell debris, and was provided for use as a sample for the subsequent ELISA. In this procedure, the purification of the chimeric antibody from the COS-7 cell culture supernatant was performed using AffiGel Protein A MAPSII Kit (Bio Rad) in accordance with the instructions included in the kit.

#### (5) ELISA

##### (i) Determination of antibody concentration

**[0207]** An ELISA plate for determining antibody concentration was prepared as follows. Each well of a 96-well ELISA plate (Maxisorp, NUNC) was coated with 100 µl of a coating buffer (0.1 M NaHCO<sub>3</sub>, 0.02% NaN<sub>3</sub>) supplemented with 1 µg/ml of goat anti-human IgG antibody (TAGO), and then blocked with 200 µl of a dilution buffer [50 mM Tris-HCl, 1 mM MgCl<sub>2</sub>, 0.1 M NaCl, 0.05% Tween 20, 0.02% NaN<sub>3</sub>, 1% bovine serum albumin (BSA); pH 7.2]. Each well of the plate was added with each of the serial dilutions of the COS-7 cell culture supernatant in which each of the chimeric antibodies had been expressed, or added with each of the serial dilutions of each of the chimeric antibodies per se in a purified form. The plate was incubated at room temperature for 1 hour and washed with PBS-Tween 20. Each well of the plate was then added with 100 µl of a solution of alkaline phosphatase-conjugated goat anti-human IgG antibodies (TAGO). After the plate was incubated at room temperature for 1 hour and washed with PBS-Tween 20, each well was added with 1mg/ml of a substrate solution ("Sigma 104", p-nitrophenylphosphoric acid, SIGMA). The solution was measured on its absorbance at 405 nm using Microplate Reader (Bio Rad) to determine the antibody concentration. In this determination, Hu IgG1 λ Purified (The Binding Site) was used as the standard substance.

##### (ii) Determination of antigen-binding ability

**[0208]** An ELISA plate for the determination of antigen-binding ability was prepared as follows. Each well of a 96-well ELISA plate was coated with 100 µl of a coating buffer supplemented with 1 µg/ml of human PTHrP (1-34) (Peptide Research Institute), and then blocked with 200 µl of a dilution buffer. Each well was added with each of the serial dilutions of the COS-7 cell culture supernatant in which each of the chimeric antibodies had been expressed, or added with each of the serial dilutions of each of the chimeric antibodies per se in a purified form. After the plate was incubated at room temperature and washed with PBS-Tween 20, each well of the plate was added with 100 µl of a solution of alkaline phosphatase-conjugated goat anti-human IgG antibodies (TAGO). After the plate was incubated at room temperature and washed with PBS-Tween 20, each well of the plate was added with 1 mg/ml of a substrate solution ("Sigma

104", p-nitrophenylphosphoric acid, SIGMA). The solution was measured on its absorbance at 405 nm using Microplate Reader (Bio Rad).

[0209] As a result, it was found that the chimeric antibodies had an ability to bind to human PTHrP (1-34) and the cloned mouse antibody V-regions had the correct structures (FIG. 5). It was also found that there was no difference in the ability to bind to PTHrP (1-34) between the chimeric antibody with L-chain  $\lambda$  chain C-region and the chimeric antibody with L-chain  $\kappa$  chain C-region. Therefore, the humanized antibody L-chain  $\lambda$  chain was used for construction of the L-chain C-region of the humanized antibody.

(6) Establishment of CHO cell line capable of stable production of chimeric antibodies

[0210] To establish a cell line capable of producing the chimeric antibodies stably, the above-prepared expression plasmids were introduced into CHO cells (DXB11).

[0211] For the establishment of a cell line capable of producing the chimeric antibodies stably, either of the following combinations of the expression plasmids for CHO cell was used: MBC1HcDNA/pCHO1 and MBC1L( $\lambda$ )/neo; and MBC1HcDNA/pCHO1 and MBC1L( $\kappa$ )/neo. A CHO cell was co-transfected with the plasmids by electroporation using Gene Pulser (Bio Rad) as follows. The expression vectors were separately cleaved with a restriction enzyme PvuI to give linear DNAs. The resulting DNAs were extracted with phenol and chloroform and collected by precipitation with ethanol. The plasmid DNAs thus prepared were subjected to electroporation. That is, each of the plasmid DNAs (10  $\mu$ g each) was added to 0.8 ml of a cell suspension of CHO cells in PBS(-) ( $1 \times 10^7$  cells/ml). The resulting solution was applied with pulses at an electrostatic capacity of 1,500V and 25  $\mu$ F. After 10 min. of recovery period at room temperature, the electroporated cells were suspended in MEM- $\alpha$  medium (GIBCO) containing 10% fetal calf serum (GIBCO). The resulting suspension was cultured using three 96-well plates (Falcon) in a CO<sub>2</sub> incubator. On the day following the culturing being started, the medium was replaced by a selective medium [ribonucleoside- or deoxyribonucleoside-free MEM- $\alpha$  medium (GIBCO) containing 10% fetal calf serum (GIBCO) and 500 mg/ml of GENETICIN (G418Sulfate; GIBCO)]. From the culture medium, cells into which the antibody gene was introduced were selected. The selective medium is replaced by a fresh one. About two weeks after the medium replacement, the cells were observed under a microscope. When a satisfactory cell growth was observed, the amount of the antibodies produced was determined by ELISA as set forth above. Among the cells, those cells which produced a larger amount of antibodies were screened.

[0212] Then, the culturing of the established cell line capable of stable production of the antibodies was scaled up in a roller bottle using ribonucleoside- or deoxyribonucleoside-free MEM medium containing 2% Ultra Low IgG fetal calf serum. On day 3 and day 4 of the culturing, the culture supernatant was collected and then filtered on a 0.2-  $\mu$ m filter (Millipore) to remove cell debris therefrom.

[0213] Purification of the chimeric antibodies from the CHO cell culture supernatant was performed using POROS Protein A Column (PerSeptive Biosystems) on ConSep LC100 (Millipore) in accordance with the instructions included in the kit. The purified chimeric antibodies were provided for use as samples for the determination of neutralizing activity and for the examination of therapeutic efficacy in hypercalcemic model animals. The concentration and the antigen-binding activity of the purified chimeric antibodies were determined using the same ELISA system as set forth above.

[REFERENCE EXAMPLE 4] Construction of humanized antibody

(1) Construction of humanized antibody H-chain

(i) Construction of humanized H-chain V-region

[0214] A humanized #23-57-137-1 antibody H-chain was produced by CDR-grafting technique by means of PCR method. For the production of a humanized #23-57-137-1 antibody H-chain (version "a") having FRs derived from human antibody S31679 (NBRF-PDB; Cuisinier, A. M. et al., Eur. J. Immunol., 23, 110-118, 1993), the following six PCR primers were used: CDR-grafting primers: MBC1HGP1 (SEQ ID NO: 23) and MBC1HGP3 (SEQ ID NO: 24) (both containing a sense DNA sequence) and MBC1HGP2 (SEQ ID NO: 25) and MBC1HGP4 (SEQ ID NO: 26) (both containing an antisense DNA sequence), all of which containing a 15-21 bp complementary sequence on both terminal ends thereof; and external primers: MBC1HVS1 (SEQ ID NO: 27) and MBC1HVR1 (SEQ ID NO: 28) having a homology to the CDR-grafting primers MBC1HGP1 and MBC1HGP4, respectively.

[0215] The CDR-grafting primers MBC1HGP1, MBC1HGP2, MBC1HGP3 and MBC1HGP4 were separated on an urea-denatured polyacrylamide gel (Molecular Cloning: A Laboratory Manual, Sambrook, et al., Cold Spring Harbor Laboratory Press, 1989), and extracted therefrom by crush-and-soak method (Molecular Cloning: A Laboratory Manual, Sambrook, et al., Cold Spring Harbor Laboratory Press, 1989) in the following manner.

[0216] Each of the CDR-grafting primers (1 nmole) was separated on a 6% denatured polyacrylamide gel to give DNA fragments. From the resulting DNA fragments, a DNA fragment having a desired length was identified on a silica

gel thin plate by irradiation of UV ray and then collected therefrom by crush-and-soak method. The resulting DNA was dissolved in 20  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA. The PCR reaction was performed using TaKaRa Ex Taq (Takara Shuzo Co., Ltd.). The PCR reaction solution (100  $\mu$ l) comprised 1  $\mu$ l of each of the above-mentioned CDR-grafting primers MBC1HGP1, MBC1HGP2, MBC1HGP3 and MBC1HGP4, 0.25 mM dNTPs and 2.5U of TaKaRa Ex Taq in the buffer. The PCR reaction was run for 5 cycles under the conditions: 94° C for 1 min., 55° C for 1 min. and 72° C for 1 min. The resulting reaction solution was added with the external primers MBC1HVS1 and MBC1HVR1 (50 pmoles each). Using this reaction mixture, the PCR reaction was run for additional 30 cycles under the same conditions. The DNA fragment thus amplified was separated by agarose gel electrophoresis on a 4% Nu Sieve GTG agarose (FMC Bio. Products).

**[0217]** An agarose segment containing a DNA fragment of 421 bp was excised, and the DNA fragment was purified therefrom using GENECLANII Kit (BIO101) in accordance with the instructions included in the kit. The DNA fragment thus purified was precipitated with ethanol and then dissolved in 20  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA. The resulting PCR reaction mixture was used for subcloning of the DNA fragment into plasmid pUC19 that had been digested with BamHI and HindIII, and subsequently the nucleotide sequence of the resulting plasmid was determined. A plasmid having the correct nucleotide sequence was designated "hMBC1Hv/pUC19".

#### (ii) Construction of H-chain V-region of Humanized H-chain cDNA

**[0218]** To ligate to cDNA for humanized H-chain C-region C  $\gamma$  1, the DNA for the humanized H-chain V-region constructed in the above step was modified by PCR method. For the PCR method, a backward primer MBC1HVS2 was designed to hybridize to the sequence encoding the 5' region of the leader sequence for the V-region and to have a Kozak consensus sequence (Kozak et al., J. Mol. Biol. 196, 947-950, 1987) and HindIII- and EcoRI-recognition sequences; and a forward primer MBC1HVR2 was designed to hybridize to both the DNA sequence encoding the 3' region of the J region and the DNA sequence encoding the 5' region of the C-region and to have ApaI- and SmaI-recognition sequences.

**[0219]** The PCR reaction was performed using TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) and a buffer appended thereto. The PCR reaction solution comprised 0.4  $\mu$ g of hMBC1Hv/pUC19 as a DNA template, 50 pmoles of each of MBC1HVS2 and MBC1HVR2 as primers, 2.5U of TaKaRa Ex Taq and 0.25 mM dNTPs in the buffer. The PCR reaction was run for 30 cycles under the conditions: 94° C for 1 min., 55° C for 1 min. and 72° C for 1 min. The DNA fragment thus amplified was separated by agarose gel electrophoresis on a 3% Nu Sieve GTG agarose (FMC Bio. Products).

**[0220]** A gel segment containing a DNA fragment of 456 bp was excised, and the DNA fragment was purified therefrom using GENECLANII Kit (BIO101) in accordance with the instructions included in the kit. The DNA fragment thus purified was precipitated with ethanol and then dissolved in 20  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA. The PCR reaction solution thus obtained was used for subcloning of the DNA fragment into plasmid pUC19 that had been digested with EcoRI and SmaI, and then the resulting plasmid was sequenced. As a result, a plasmid was obtained which contained a DNA encoding mouse H-chain V-region derived from hybridoma #23-57-137-1 and also contained EcoRI- and HindIII-recognition sequences and a Kozak sequence on the 5' region and ApaI- and SmaI-recognition sequences on the 3' region, which was designated "hMBC1Hv/pUC19".

#### (2) Construction of expression vector for humanized antibody H-chain

**[0221]** Plasmid RVh-PM1f-cDNA carrying a cDNA sequence for hPM1 antibody H-chain was digested with ApaI and BamHI to give a DNA fragment containing a DNA fragment containing a DNA encoding the H-chain C-region. The DNA fragment was introduced into plasmid hMBC1Hv/pUC19 that had been digested with ApaI and BamHI. The obtained plasmid was designated "hMBC1HcDNA/pUC19". This plasmid contained both a DNA encoding the humanized #23-57-137-1 antibody H-chain V-region and a DNA encoding the human H-chain C-region C  $\gamma$  1 and had EcoRI- and HindIII-recognition sequences on the 5' region and a BamHI-recognition sequence on the 3' region. The nucleotide sequence and the corresponding amino acid sequence of the humanized H-chain version "a" carried on the plasmid hMBC1HcDNA/pUC19 are shown in SEQ ID NO: 58 and SEQ ID NO: 56, respectively.

**[0222]** The plasmid hMBC1HcDNA/pUC19 was digested with EcoRI and BamHI to give a DNA fragment containing a DNA encoding the H-chain. The DNA fragment was introduced into expression plasmid pCOS1 that had been digested with EcoRI and BamHI. As a result, an expression plasmid for a humanized antibody was obtained, which was designated "hMBC1HcDNA/pCOS1".

**[0223]** To produce a plasmid used for expression in a CHO cell, plasmid hMBC1HcDNA/pUC19 was digested with EcoRI and BamHI to give a DNA fragment containing a DNA encoding the H-chain. The DNA fragment was introduced into expression vector pCHO1 that had been digested with EcoRI and BamHI. As a result, an expression plasmid for the humanized antibody was obtained, which was designated "hMBC1HcDNA/pCHO1".

## (3) Construction of L-chain hybrid V-region

## (i) Preparation of FR1,2/FR3,4 hybrid antibody

- [0224]** A gene for the FR hybrid L-chain having both FRs from a humanized antibody and FRs from a mouse (chimeric) antibody was constructed, and evaluated each region for the humanization. In this step, a hybrid antibody having FR1 and FR2 both derived from a human antibody and FR3 and FR4 both derived from a mouse antibody was prepared by utilizing the AflIII restriction site located on CDR2.
- [0225]** Plasmids MBC1L( $\lambda$ )/neo and hMBC1L( $\gamma$ )/neo (10  $\mu$ g each) were separately digested in 100  $\mu$ l of a reaction solution containing 10 mM Tris-HCl (pH 7.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 50 mM NaCl, 0.01% (w/v) of BSA and 10 U of AflIII (Takara Shuzo Co., Ltd.) at 37 °C for 1 hour. The reaction solutions were subjected to electrophoresis on a 2% low-melting agarose gel, thereby giving DNA fragments of 6282 bp (referred to as "c1") and 1022 bp (referred to as "c2") from the plasmid MBC1L( $\lambda$ )/neo or DNA fragments of 6282 bp (referred to as "h1") and 1022 bp (referred to as "h2") from the plasmid hMBC1L( $\lambda$ )/neo. These DNA fragments were collected and purified from the gels using GENE-CLEANII Kit (BIO101).
- [0226]** Each of the c1 and h1 fragments (1  $\mu$ g each) was BAP-treated. The DNA fragment was extracted with phenol and chloroform, collected by ethanol precipitation, and then dissolved in 10  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.
- [0227]** The BAP-treated c1 and h1 DNA fragments (1  $\mu$ l each) were ligated to the h2 and c2 DNA fragments (4  $\mu$ l 1 each), respectively, (at 4° C overnight). Each of the ligation products was introduced into a competent cell of E. coli, JM109, to form a transformant. The transformant was cultured in 2 ml of 2xYT medium containing 50  $\mu$ g/ml of ampicillin. From the cell fraction, the plasmid was purified using QIAprep Spin Plasmid Kit (QIAGEN).
- [0228]** The purified plasmid was digested in 20  $\mu$ l of a reaction solution containing 10 mM Tris-HCl (pH 7.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, and either 2U of ApaLI (Takara Shuzo Co., Ltd.) or 8U of BamHI (Takara Shuzo Co., Ltd.) and HindIII (Takara Shuzo Co., Ltd.) at 37° C for 1 hour. It was expected that if the c1-h2 was ligated correctly, this digestion reaction would give fragments of 5560/1246/498 bp (by the ApaLI digestion) or fragments of 7134/269 bp (by the BamHI/HindIII digestion). Based on this expectation, the desired plasmids were identified.
- [0229]** The expression vector encoding the human FR1,2/mouse FR3,4 hybrid antibody L-chain was designated "h/mMBC1L( $\lambda$ )/neo". On the other hand, since a clone for the h1-c1 could not be obtained, recombination on a pUC vector was performed and then the resulting recombinant product was cloned into a HEF vector. In this procedure, plasmid hMBC1La  $\lambda$  /pUC19, which contained DNA encoding a humanized antibody L-chain V-region without any amino acid replacements, and plasmid hMBC1Ld  $\lambda$  /pUC19, which contained a DNA encoding a humanized antibody L-chain V-region with an amino acid replacement at the 91-position amino acid tyrosine in FR3 (i.e., the 87th amino acid in accordance with The Kabat's prescription) by isoleucine, were used as templates.
- [0230]** Plasmids MBC1L( $\lambda$ )/pUC19, hMBC1La $\lambda$ /pUC19 and hMBC1Ld $\lambda$ /pUC19 (10  $\mu$ l each) were separately digested in 30  $\mu$ l of a reaction solution containing 10 mM Tris-HCl (pH 7.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 50 mM NaCl, 0.01 % (w/v) of BSA, 16U of HindIII and 4U of AflIII at 37° C for 1 hour. The reaction solutions were separately subjected to electrophoresis on a 2% low-melting agarose gel, thereby giving a DNA fragment of 215 bp from plasmid MBC1L( $\lambda$ )/pUC19 (referred to as "c2'") and a DNA fragment of 3218 bp from each of plasmids hMBC1La $\lambda$ /pUC19 and hMBC1Ld $\lambda$ /pUC19 (referred to as "ha1'" and "hd1'", respectively). These DNA fragments were collected and purified using GENE-CLEANII Kit (BIO101).
- [0231]** Each of the ha1' and hd1' fragments was ligated to the c2' fragment and then introduced into a competent cell of E. coli, JM109, to form a transformant. The transformant was cultured in 2 ml of 2xYT medium containing 50  $\mu$ g/ml of ampicillin. From the cell fraction, the plasmid was purified using QIAprep Spin Plasmid Kit (QIAGEN). The plasmids thus prepared were designated "m/hMBC1La $\lambda$ /pUC19" for the ha1' fragment-containing plasmid and "m/hMBC1Ld $\lambda$ /pUC19" for the hd1' fragment-containing plasmid.
- [0232]** Each of the plasmids m/hMBC1La $\lambda$ /pUC19 and m/hMBC1Ld $\lambda$ /pUC19 was digested with EcoRI. The DNA fragment of 743 bp was electrophoresed on a 2% low-melting agarose gel, and then collected and purified therefrom using GENE-CLEANII Kit (BIO101). The resulting DNA fragment was dissolved in 20  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.
- [0233]** Each of the DNA fragments (4  $\mu$ l each) was ligated to the above-obtained BAP-treated HEF vector (1  $\mu$ l). The ligation product was introduced into a competent cell of E. coli, JM109, to form a transformant. The transformant was cultured in 2 ml of 2xYT medium containing 50  $\mu$ g/ml of ampicillin. From the cell fraction, the plasmid was purified using QIAprep Spin Plasmid Kit (QIAGEN).
- [0234]** Each of the purified plasmids was digested in 20  $\mu$ l of a reaction solution containing 20 mM Tris-HCl (pH 8.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 100 mM KCl, 8U of HindIII (Takara Shuzo Co., Ltd.) and 2U of PvuI (Takara Shuzo Co., Ltd.) at 37 °C for 1 hour. It was expected that if the DNA fragment was inserted in the plasmid in a correct orientation, this digestion would give digestion fragments of 5104/2195 bp, whereas if the DNA fragment is inserted in the plasmid

in the reverse orientation, this digestion would give digestion fragments of 4378/2926 bp. The plasmid DNA was identified based on the expectation. The plasmids thus obtained were expression vectors encoding mouse FR1,2/human FR3,4 hybrid antibody L-chain, which were designated expression vectors "m/hMBC1La $\lambda$ /neo" and "m/hMBC1Ld $\lambda$ /neo", respectively.

#### (ii) Preparation of FR1/FR2 hybrid antibody

**[0235]** An FR1/FR2 hybrid antibody was prepared in the same manner as set forth above utilizing a SnaBI restriction site located on CDR1.

**[0236]** Plasmids MBC1L( $\lambda$ )/neo and h/mMBC1L( $\lambda$ )/neo (10  $\mu$ g each) were separately digested in 20  $\mu$ l of a reaction solution containing 10 mM Tris-HCl (pH 7.9), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 50 mM NaCl, 0.01% (w/v) of BSA and 6U of SnaBI (Takara Shuzo Co., Ltd.) at 37 ° C for 1 hour. The resulting reaction solutions were further digested in 50  $\mu$ l of a reaction solution containing 20 mM Tris-HCl (pH 8.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT, 100 mM KCl, 0.01 % (w/v) of BSA and 6U of PvuI at 37° C for 1 hour.

**[0237]** The resulting reaction solutions were separately subjected to electrophoresis on a 1.5% low-melting agarose gel, thereby giving DNA fragments of 4955 bp (m1) and 2349 bp (m2) from the plasmid MBC1L( $\lambda$ )/neo and DNA fragments of 4955 bp (hm1) and 2349 bp (hm2) from the plasmid h/mMBC1L( $\lambda$ )/neo. These DNA fragments were collected and purified from the gels using GENECLANII Kit (BIO101). Each of the DNA fragments obtained was dissolved in 40  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

**[0238]** The m1 and hm1 fragments (1  $\mu$ l each) were ligated to the hm2 and m2 fragments (4  $\mu$ l each), respectively. Each of the resulting ligation products was introduced into a competent cell of E. coli, JM109, to form a transformant. The transformant obtained was cultured in 2 ml of 2xYT medium containing 50  $\mu$ g/ml of ampicillin. From the cell fraction, the plasmid was purified using QIAprep Spin Plasmid Kit QIAGEN).

**[0239]** Each of the purified plasmids was digested in 20  $\mu$ l of a reaction solution containing 10 mM Tris-HCl (pH 7.5), 10 mM MgCl<sub>2</sub>, 1 mM DTT and either 8U of Apal (Takara Shuzo Co., Ltd.) or 2U of ApaI (Takara Shuzo Co., Ltd.) at 37° C for 1 hour.

**[0240]** It was expected that if the fragments were ligated correctly, the digestion reaction would give a fragment of 7304 bp (by the Apal digestion) or fragments of 5560/1246/498 bp (by the ApaI digestion) for m1-hm2, and would give fragments of 6538/766 bp (by the Apal digestion) or fragments of 3535/2025/1246/498 bp (by the ApaI digestion) for hm1-m2. Based on this expectation, the plasmids were identified. As a result, an expression vector encoding a human FR1/mouse FR2,3,4 hybrid antibody L-chain (designated "hmmMBC1L( $\lambda$ )/neo") and an expression vector encoding a mouse FR1/human FR2/mouse FR3,4 hybrid antibody L-chain (designated "mhmMBC1L( $\lambda$ )/neo") were obtained.

#### (4) Construction of humanized antibody L-chain

**[0241]** A humanized #23-57-137-1 antibody L-chain was prepared by CDR-grafting technique by means of PCR method. For the preparation of a humanized #23-57-137-1 antibody L-chain (version "a") that contained FR1, FR2 and FR3 derived from human antibody HSU03868 (GEN-BANK, Deftos M. et al., Scand. J. Immunol., 39, 95-103, 1994) and FR4 derived from human antibody 525755 (NBRF-PDB), six PCR primers were used.

**[0242]** The six primers were as follows: CDR-grafting primers MBC1LGP1 (SEQ ID NO: 29) and MBC1LGP3 (SEQ ID NO: 30), both having a sense DNA sequence, CDR-grafting primers MBC1LGP2 (SEQ ID NO: 31) and MBC1LGP4 (SEQ ID NO: 32), both having an antisense DNA sequence, all of which had a 15-21 bp complementary sequence on the both terminal ends; and external primers MBC1LVS1 (SEQ ID NO: 33) and MBC1LVR1 (SEQ ID NO: 34) having a homology to the CDR-grafting primers MBC1LGP1 and MBC1LGP4, respectively.

**[0243]** The CDR-grafting primers MBC1LGP1, MBC1LGP2, MBC1LGP3 and MBC1LGP4 were separated on a urea-denatured polyacrylamide gel (Molecular Cloning: A Laboratory Manual, Sambrook et al., Cold Spring Harbor Laboratory Press, 1989) and extracted therefrom by crush-and-soak method (Molecular Cloning: A Laboratory Manual, Sambrook et al., Cold Spring Harbor Laboratory Press, 1989).

**[0244]** Each of the CDR-grafting primers (1 nmole each) was separated on a 6% denatured polyacrylamide gel. The identification of the DNA fragment of a desired length was performed on a silica gel thin plate by irradiation of UV ray. The desired DNA fragment was collected from the gel by crush-and-soak method. The collected DNA fragment was dissolved in 20  $\mu$ l of a solution containing 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA.

**[0245]** The PCR reaction was performed using TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) and a buffer appended thereto. The PCR reaction solution comprised (per 100  $\mu$ l) 1  $\mu$ l of each of the CDR-grafting primers MBC1LGP1, MBC1LGP2, MBC1LGP3 and MBC1LGP4, 0.25 mM dNTPs, 2.5U of TaKaRa Ex Taq in the buffer. The PCR reaction was run for 5 cycles under the conditions: 94° C for 1 min., 55° C for 1 min. and 72° C for 1 min. The resulting reaction mixture was added with 50 pmoles of each of the external primers MBC1LVS1 and MBC1LVR1. Using this reaction

mixture, the PCR reaction was run for additional 30 cycles under the same conditions. The DNA fragment thus amplified was separated by agarose gel electrophoresis on a 3% Nu Sieve GTG agarose (FMC Bio. Products).

[0246] An agarose segment containing a DNA fragment of 421 bp was excised, and the DNA fragment was purified therefrom using GENECLANII Kit (BIO101) in accordance with the instructions included in the kit. The PCR reaction mixture thus obtained was used for subcloning of the DNA fragment into plasmid pUC19 that had been digested with BamHI and HindIII. The resulting plasmid was sequenced. The plasmid thus prepared was designated "hMBCL/pUC19". In this plasmid, however, the 104-position amino acid (corresponding to the 96th amino acid in accordance with the Kabat's prescription) of CDR4 was replaced by arginine. For the correction of this amino acid to tyrosine, a correction primer MBC1LGP10R (SEQ ID NO: 35) was designed and synthesized. The PCR reaction was performed using TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) and a buffer appended thereto. The PCR reaction solution comprised (per 100  $\mu$ l) 0.6  $\mu$ g of the plasmid hMBCL/pUC19 as a template DNA, 50 pmoles of each of the primers MBC1LVS1 and MBC1LGP10R, 2.5U of TaKaRa Ex Taq (Takara Shuzo Co., Ltd.) and 0.25 mM dNTPs in the buffer, over which mineral oil (50  $\mu$ l) was layered. The PCR reaction was run for 30 cycles under the conditions: 94° C for 1 min., 55° C for 1 min. and 72° C for 1 min. The DNA fragment thus amplified was separated by agarose gel electrophoresis on a 3% Nu Sieve GTG agarose (FMC Bio. Products).

[0247] A gel segment containing a DNA fragment of 421 bp was excised, and the DNA fragment was purified therefrom using GENECLANII Kit (BIO101) in accordance with the instructions included in the kit. The PCR reaction mixture thus prepared was used for subcloning of the DNA fragment into plasmid pUC19 that had been digested with BamHI and HindIII.

[0248] The plasmid was sequenced using M13 Primer M4 and M13 Primer RV. As a result, it was confirmed that the plasmid had the correct sequence. The plasmid was then digested with HindIII and BlnI, and a DNA fragment of 416 bp was separated by electrophoresis on a 1% agarose gel. The DNA fragment was purified using GENECLANII Kit (BIO101) in accordance with the instructions included in the kit, and then introduced into plasmid C $\lambda$ /pUC19 that had been digested with HindIII and BlnI. The resulting plasmid was designated "hMBC1La $\lambda$ /pUC19". This plasmid was digested with EcoRI to give a DNA fragment encoding humanized L-chain. The DNA fragment was introduced into plasmid pCOS1 so that the initiation codon for the humanized L-chain was located downstream to the EF1 $\alpha$  promoter. The plasmid thus obtained was designated "hMBC1La $\lambda$ /pCOS1". The DNA sequence (including the corresponding amino acid sequence) of the humanized L-chain version "a" is shown in SEQ ID NO: 66. The amino acid sequence of the version "a" is also shown in SEQ ID NO: 47.

[0249] A humanized L-chain version "b" was prepared using mutagenesis by PCR method. The version "b" was designed such that the 43-position amino acid glycine (corresponding to the 43th amino acid in accordance with the Kabat's prescription) was replaced by proline and the 49-position amino acid lysine (corresponding to the 49th amino acid in accordance with the Kabat's prescription) by aspartic acid in the version "a". The PCR reaction was performed using plasmid hMBC1La $\lambda$ /pUC19 as a template and a mutagenic primer MBC1LGP5R (SEQ ID NO: 36) and a primer MBC1LVS1. The DNA fragment obtained was digested with BamHI and HindIII, and the digestion fragment was subcloned into the BamHI-HindIII site of pUC19. After sequencing, the plasmid was digested with HindIII and AflII, and the resulting digestion fragment was ligated to plasmid hMBC1La $\lambda$ /pUC19 that had been digested with HindIII and AflII.

[0250] The plasmid thus obtained was designated "hMBC1Lb $\lambda$ /pUC19". This plasmid was digested with EcoRI to give a DNA fragment containing a DNA encoding the humanized L-chain. The DNA fragment was introduced into plasmid pCOS1 such that the initiation codon for the humanized L-chain was located downstream to the EF1 $\alpha$  promoter. The plasmid thus obtained was designated "hMBC1Lb $\lambda$ /pCOS1".

[0251] A humanized L-chain version "c" was prepared using mutagenesis by PCR method. The version "c" was designed such that the 84-position amino acid serine (corresponding to the 80th amino acid in accordance with the Kabat's prescription) was replaced by proline. The PCR reaction was performed using plasmid hMBC1La $\lambda$ /pUC19 as a template and a mutagenic primer MBC1LGP6S (SEQ ID NO: 37) and a primer M13 Primer RV. The DNA fragment obtained was digested with BamHI and HindIII and then subcloned into pUC19 that had been digested with BamHI and HindIII.

[0252] After sequencing, the plasmid was digested with BstPI and Aor51HI, and the resulting DNA fragment was ligated to plasmid hMBC1La $\lambda$ /pUC19 that had been digested with BstPI and Aor51HI. The plasmid thus obtained was designated "hMBC1Lc $\lambda$ /pUC19". This plasmid was digested with EcoRI to give a DNA fragment containing a DNA encoding the humanized L-chain. The fragment was introduced into the EcoRI site of plasmid pCOS1 such that the initiation codon for the humanized L-chain was located downstream to the EF1 $\alpha$  promoter. The plasmid thus obtained was designated "hMBC1Lc $\lambda$ /pCOS1".

[0253] Humanized L-chain versions "d", "e" and "f" were also prepared using mutagenesis by PCR method. The versions "d", "e" and "f" were designed such that the 91-position amino acid tyrosine (corresponding to the 87th amino acid in accordance with the Kabat's prescription) was replaced by isoleucine in the versions "a", "b" and "c", respectively. For each of the versions "d", "e" and "f", a PCR reaction was performed using each of plasmid hMBC1La $\lambda$ /pCOS1 (for version "d"), hMBC1Lb $\lambda$ /pCOS1 (for version "e") and hMBC1Lc $\lambda$ /pCOS1 (for version "f"), respectively, as a template,

a mutagenic primer MBC1LGP11R (SEQ ID NO: 38) and a primer M-S1 (SEQ ID NO: 44). The DNA fragment thus obtained was digested with BamHI and HindIII and then subcloned into pUC19 that had been digested with BamHI and HindIII. After sequencing, the plasmid was digested with HindIII and BlnI, and the resulting digestion fragment was ligated to plasmid Cλ/pUC19 that had been digested with HindIII and BlnI.

**[0254]** The plasmids thus obtained were respectively designated "hMBC1Ldλ/pUC19" (for version "d"), "hMBC1Leλ/pUC19" (for version "e") and "hMBC1Lfλ/pUC19" (for version "f"). Each of these plasmids was digested with EcoRI to give a DNA fragment containing a DNA encoding the humanized L-chain. The DNA fragment was introduced into the EcoRI site of plasmid pCOS1 such that the initiation codon for the humanized L-chain was located downstream to the EF1 α promoter of the plasmid. The plasmids thus obtained were respectively designated "hMBC1Ldλ/pCOS1" (for version "d"), "hMBC1Leλ/pCOS1" (for version "e") and "hMBC1Lfλ/pCOS1" (for version "f").

**[0255]** Humanized L-chain versions "g" and "h" were also prepared using mutagenesis by PCR method. The versions "g" and "h" were designed such that the 36-position amino acid histidine (corresponding to the 36th amino acid in accordance with the Kabat's prescription) was replaced by tyrosine in the versions "a" and "d", respectively. The PCR reaction was performed using a mutagenic primer MBC1LGP9R (SEQ ID NO: 39), M13 Primer RV and plasmid hMBC1Laλ/pUC19 as a template. An additional PCR was performed using the PCR product thus obtained and M13 Primer M4 as primers and plasmid hMBC1Laλ/pUC19 as a template. The DNA fragment obtained was digested with HindIII and BlnI and then subcloned into plasmid Cλ/pUC19 that had been digested with HindIII and BlnI. Using this plasmid as a template, a PCR reaction was performed using primers MBC1LGP13R (SEQ ID NO: 40) and MBC1LVS1. The PCR fragment obtained was digested with ApaI and HindIII and then introduced into either of plasmids hMBC1Laλ/pUC19 and hMBC1Ldλ/pUC19 that had been digested with ApaI and HindIII. The plasmids obtained were sequenced. Plasmids that were confirmed to contain the correct sequence were designated "hMBC1Lgλ/pUC19" (for version "g") and "hMBC1Lhλ/pUC19" (for version "h"). Each of these plasmids was digested with EcoRI to give a DNA fragment containing a DNA encoding the humanized L-chain. The DNA fragment was introduced into the EcoRI site of plasmid pCOS1 such that the initiation codon for the humanized L-chain was located downstream to the EF1 α promoter. The plasmids thus obtained were respectively designated "hMBC1Lgλ/pCOS1" (for version "g") and "hMBC1Lhλ/pCOS1" (for version "h").

**[0256]** Humanized L-chain versions "i", "j", "k", "l", "m", "n" and "o" were also prepared using mutagenesis by PCR method. The PCR reaction was performed using plasmid hMBC1Laλ/pUC19 as a template and a mutagenic primer MBC1LGP14S (SEQ ID NO: 41) and a primer V1RV (λ) (SEQ ID NO: 43). The resulting DNA fragment was digested with ApaI and BlnI and then subcloned into plasmid hMBC1Lgλ/pUC19 that had been digested with ApaI and BlnI. The plasmid obtained was sequenced, and the clone into which the mutation for each version was introduced was selected. The plasmid thus obtained was designated "hMBC1Lxλ/pUC19 (x=i, j, k, l, m, n or o)". This plasmid was digested with EcoRI to give a DNA fragment containing a DNA encoding the humanized L-chain. The DNA fragment was introduced into the EcoRI site of plasmid pCOS1 such that the initiation codon for the humanized L-chain was located downstream to the EF1 α promoter. The plasmid thus obtained was designated "hMBC1Lxλ/pCOS1" (x = i, j, k, l, m, n or o). The DNA sequences (including the corresponding amino acid sequences) of the versions "j", "l", "m" and "o" are shown in SEQ ID NOs: 67, 68, 69 and 70, respectively. The amino acid sequences of these versions are also shown in SEQ ID Nos: 48, 49, 50 and 51, respectively.

**[0257]** Humanized L-chain versions "p", "q", "r", "s" and "t" were designed such that the 87-position amino acid (tyrosine) was replaced by isoleucine in the versions "i", "j", "m", "l" and "o", respectively. These versions were prepared utilizing an Aor51HI restriction site on FR3 and replacing that site of each of the versions "i", "j", "m", "l" or "o" by that site of the version "h". That is, an Aor51HI restriction fragment (514 bp) containing CDR3, a part of FR3 and the entire FR4 were removed from an expression plasmid hMBC1Lxλ/pCOS1 (x = i, j, m, l or o). To the removed site, an Aor51HI restriction fragment (514 bp) in the expression plasmid hMBC1Lhλ/pCOS, which containing CDR3 and a part of FR3 and the entire FR4, was ligated, so that the 91-position amino acid tyrosine (corresponding to the 87th amino acid in accordance with the Kabat's prescription) was replaced by isoleucine. The resulting plasmid was sequenced. A clone of each of the versions "i", "j", "m" "l" and "o" in which 91-position amino acid tyrosine (corresponding to the 87th amino acid in accordance with the Kabat's prescription) was replaced by isoleucine was selected. These modified versions respectively corresponding to the versions "i", "j", "m" "l" and "o" were designated versions "p", "q", "s", "r" and "t", respectively. The obtained plasmid was designated "hMBC1Lxλ/pCOS1 (x = p, q, s, r or t)". The DNA sequences (including the corresponding amino acids) of the versions "q", "r", "s" and "t" are shown in SEQ ID Nos: 71, 72, 73 and 74, respectively. The amino acid sequences of these versions are also shown in SEQ ID Nos: 52, 53, 54 and 55, respectively.

**[0258]** Plasmid hMBC1Lqλ/pCOS1 was digested with HindIII and EcoRI and then subcloned into plasmid pUC19 that had been digested with HindIII and EcoRI. The plasmid thus obtained was designated "hMBC1Lqλ/pUC19".

**[0259]** The positions of the replaced amino acids in the individual versions of the humanized L-chain are shown in Table 8 below.



Table 8

Positions of replaced amino acid in sequence listings (amino acid numbering in accordance with the Kabat's prescription)							
Versions	36	43	45	47	49	80	87
a							
b		P			D		
c						P	
d							I
e		P			D		I
f						P	I
g	Y						
h	Y						I
i	Y		K				
j	Y		K		D		
k	Y		K	V			
l	Y		K	V	D		
m	Y				D		
n	Y			V			
o	Y			V	D		
p	Y		K				I
q	Y		K		D		I
r	Y				D		I
s	Y		K	V	D		I
t	Y			V	D		I

In the Table, capital letters represent the following amino acids: Y: tyrosine; P: proline; K: lysine; V: valine; D: aspartic acid; and I: isoleucine.

**[0260]** E. coli strains each containing plasmids hMBC1HcDNA/pUC19 and hMBC1Lqλ/pUC19 were designated "Escherichia coli JM109 (hMBC1HcDNA/pUC19)" and "Escherichia coli JM109 (hMBC1Lqλ/pUC19)", respectively, which have been deposited under the terms of Budapest Treaty at the National Institute of Bioscience and Human-Technology, Agency of Industrial Science and Technology, Japan, (1-3, Higashi 1-chome, Tsukuba-shi, Ibaraki, Japan) on August 15, 1996, under the accession No. FERM BP-5629 for Escherichia coli JM109 (hMBC1HcDNA/pUC19), and FERM BP-5630 for Escherichia coli JM109 (hMBC1Lqλ/pUC19).

#### (5) Transfection into COS-7 cell

**[0261]** For the evaluation of the antigen-binding activity and the neutralizing activity of the hybrid antibodies and the humanized #23-57-137-1 antibodies, the above-prepared expression plasmids were expressed transiently in COS-7 cells. For the transient expression of the L-chain hybrid antibodies, each of the following combinations of plasmids were co-transfected into a COS-7 cell by electroporation using Gene Pulser (Bio Rad): hMBC1HcDNA/pCOS1 and h/mMBC1L(λ)/neo; hMBC1HcDNA/pCOS1 and m/hMBC1Laλ/neo; hMBC1HcDNA/pCOS1 and m/hMBC1Ldλ/neo; hMBC1HcDNA/pCOS1 and hmmMBC1L(λ)/neo; and hMBC1HcDNA/pCOS1 and mhmMBC1L(λ)/neo. That is, a cell suspension (0.8 ml) of COS-7 cells in PBS(-) ( $1 \times 10^7$  cells/ml) was added with each combination of the plasmid DNAs (10 μg each). The resulting solution was applied with pulses at an electrostatic capacity of 1,500V and 25 μF. After 10 min. of recovery period at room temperature, the electroporated cells were suspended in DMEM medium containing 2% Ultra Low IgG fetal calf serum (GIBCO), and then cultured using a 10-cm culture dish in a CO<sub>2</sub> incubator. After culturing for 72 hours, a culture supernatant was collected and centrifuged to remove cell debris. The solutions thus

prepared were provided for use in the ELISA below.

**[0262]** For the transient expression of the humanized #23-57-137-1 antibodies, plasmids of hMBC1HcDNA/pCOS1 and hMBC1L $\lambda$ /pCOS1 ( $x = a-t$ ) were co-transfected into a COS-7 cell using Gene Pulser (Bio Rad) in the same manner as described for the hybrid antibodies above. The culture supernatants were prepared and provided for use in the ELISA below.

**[0263]** The purification of the hybrid antibodies and the humanized antibodies from the COS-7 cell culture supernatants was performed using AffiGel Protein A MAPSII Kit (Bio Rad) in accordance with the instructions included in the kit.

#### (6) ELISA

##### (i) Determination of antibody concentration

**[0264]** An ELISA plate for determining antibody concentration was prepared as follows. Each well of a 96-well ELISA plate (Maxisorp, NUNC) was coated with 100  $\mu$ l of a coating buffer (0.1 M NaHCO<sub>3</sub>, 0.02% NaN<sub>3</sub>) containing 1  $\mu$ g/ml of goat anti-human IgG antibody (TAGO) and then blocked with 200  $\mu$ l of a dilution buffer [50 mM Tris-HCl, 1 mM MgCl<sub>2</sub>, 0.1 M NaCl, 0.05% Tween 20, 0.02% NaN<sub>3</sub>, 1% bovine serum albumin (BSA); pH 7.2]. Each of the wells was added with each of the serial dilutions of the COS cell culture supernatant in which each of the hybrid antibodies and the humanized antibodies was expressed, or added with each of the serial dilutions of each of the hybrid antibodies and humanized antibodies in a purified form. The plate was incubated at room temperature for 1 hour and washed with PBS-Tween 20. Subsequently, each of the wells was added with 100  $\mu$ l of alkaline phosphatase-conjugated goat anti-human IgG antibody (TAGO). The plate was incubated at room temperature for 1 hour and washed with PBS-Tween 20. Subsequently, each of the wells was added with 1 mg/ml of a substrate solution ("Sigma 104", p-nitrophenylphosphoric acid, SIGMA). The solution in each well was measured on its absorbance at 405 nm using Microplate Reader (Bio Rad) to determine the antibody concentration. In this determination, Hu IgG1  $\lambda$  Purified (The Binding Site) was used as the standard substance.

##### (ii) Determination of antigen-binding ability

**[0265]** An ELISA plate for determining antigen-binding ability was prepared as follows. Each well of a 96-well ELISA plate (Maxisorp, NUNC) was coated with 100  $\mu$ l of a coating buffer containing 1  $\mu$ g/ml of human PTHrP (1-34) and then blocked with 200  $\mu$ l of a dilution buffer. Subsequently, each well was added with each of the serial dilutions of the COS-7 cell culture supernatant in which each of the hybrid antibodies and humanized antibodies was expressed, or added with each of the serial dilutions of each of the hybrid antibodies and humanized antibodies in a purified form. The plate was incubated at room temperature and washed with PBS-Tween 20. Subsequently, each well was added with 100  $\mu$ l of alkaline phosphatase-conjugated goat anti-human IgG antibody (TAGO). The plate was incubated at room temperature and washed with PBS-Tween 20. Subsequently, each well was added with 1 mg/ml of a substrate solution ("Sigma 104", p-nitrophenylphosphoric acid, SIGMA). The solution was measured on its absorbance at 405 nm using Microplate Reader (Bio Rad).

#### (7) Confirmation of activities

##### (i) Evaluation of humanized H-chain

**[0266]** It was found that an antibody having both a humanized H-chain version "a" and a chimeric L-chain exhibited the same level of PTHrP-binding activity as that of a chimeric antibody. This result suggests that the version "a" achieves the humanization of the H-chain V-region in the degree enough to evaluate the humanization. Therefore, the humanized H-chain version "a" was provided for use as a humanized antibody H-chain in the following experiments.

##### (ii) Activity of hybrid antibodies

##### (ii-a) FR1,2/FR3,4 hybrid antibody

**[0267]** When the L-chain was h/mMBC1L( $\lambda$ ), no antigen-binding activity was observed. In contrast, when the L-chain was either m/hMBC1La  $\lambda$  or m/hMBC1Ld  $\lambda$ , the same level of antigen-binding activity as that of the chimeric #23-57-137-1 antibody was observed (FIG. 7). These results suggest that FR3 and FR4 have no problem as humanized antibodies but FR1 and FR2 contain amino acid residue(s) that need to be replaced.

## (ii-b) FR1/FR2 hybrid antibody

**[0268]** When the L-chain was mhmMBC1L ( $\lambda$ ), no antigen-binding activity was observed. In contrast, when the L-chain was hmmMBC1L ( $\lambda$ ), the same level of antigen-binding activity as that of the chimeric #23-57-137-1 antibody was observed (FIG. 8). These results suggest that FR1 has no problem as a humanized antibody but FR2 contains amino acid residue(s) that need to be replaced.

## (iii) Activity of humanized antibodies

**[0269]** The antigen-binding activity of the humanized antibodies having the L-chain versions "a" to "t", respectively, were determined. As a result, it was found that the humanized antibodies having the L-chain versions "j", "l", "m", "o", "q", "r", "s" and "t" exhibited the same levels of PTHrP-binding activity as that of the chimeric antibody.

## (8) Establishment of CHO cell line capable of stable production of antibody

**[0270]** For establishing a cell line capable of stable production of humanized antibodies, each of the above-prepared expression plasmids was introduced into a CHO cell (DXB11).

**[0271]** That is, the establishment of a cell line capable of stable production of a humanized antibody was performed using each of the following combinations of plasmids as expression vectors for a CHO cell; hMBC1HcDNA/pCHO1 and hMBC1Lm $\lambda$ /pCOS1; hMBC1HcDNA/pCHO1 and hMBC1Lq $\lambda$ /pCOS1; and hMBC1HcDNA/pCHO1 and hMBC1Lr $\lambda$ /pCOS1. The plasmids were co-transfected into a CHO cell by electroporation using Gene Pulser (Bio Rad). Subsequently, the expression vectors were separately cleaved with restriction enzyme PvuI to give linear DNA fragments. The resulting DNA fragments were extracted with phenol and chloroform and then precipitated with ethanol. The DNA fragments thus prepared were used in the subsequent electroporation. That is, the plasmid DNA fragments (10  $\mu$ g each) were added to 0.8 ml of a cell suspension of CHO cells in PBS(-) ( $1 \times 10^7$  cells/ml). The resulting solution was applied with pulses at an electrostatic capacity of 1,500V and 25  $\mu$ F. After 10 min. of recovery period at room temperature, the cells thus treated were suspended in MEM- $\alpha$  medium (GIBCO) containing 10% fetal calf serum (GIBCO), and then cultured in a CO<sub>2</sub> incubator using 96-well plates (Falcon). On the day following the culturing being started, the medium was replaced by ribonucleoside- or deoxyribonucleoside-free MEM- $\alpha$  selective medium containing 10% fetal calf serum (GIBCO) and 500 mg/ml of GENETICIN (G418Sulfate; GIBCO). From the culture medium, cells into which the antibody gene was introduced were selected. The culture medium was replaced by a fresh one. About two weeks after the medium replacement, the cells were observed microscopically. When a satisfactory cell growth was observed, the amount of the antibodies produced was determined by conventional ELISA for determination of antibody concentration as set forth above. Among the cells, those cells which produced a larger amount of antibodies were screened.

**[0272]** The culturing of the established cell line capable of stable production of antibodies was scaled up in a roller bottle using a ribonucleoside- or deoxyribonucleoside-free MEM- $\alpha$  medium containing 2% Ultra Low IgG fetal calf serum. On each of day 3 and day 4 of the culturing, the culture supernatant was collected and filtered on a 0.2- $\mu$ m filter (Millipore) to remove cell debris therefrom. The purification of the humanized antibodies from the culture supernatant of the CHO cells was performed using POROS Protein A Column (PerSeptive Biosystems) on ConSep LC100 (Millipore) in accordance with the appended instructions. The humanized antibodies were provided for use in the determination of neutralizing activity and examination of pharmacological efficacy in hypercalcemic model animals. The concentration and the antigen-binding activity of the purified humanized antibodies were determined by the ELISA system as set forth above.

## [REFERENCE EXAMPLE 5] Determination of neutralizing activity

**[0273]** The determination of neutralizing activity of the mouse antibodies, the chimeric antibodies and the humanized antibodies was performed using rat myeloma cell line ROS17/2.8-5 cells. The ROS17/2.8-5 cells were cultured in Ham'S F-12 medium (GIBCO) containing 10% fetal calf serum (GIBCO) in a CO<sub>2</sub> incubator. The ROS17/2.8-5 cells were seeded into each well of a 96-well plate at a density of  $10^4$  cells/100  $\mu$ l/well and cultured for one day. After the culturing was completed, the culture medium was replaced by Ham'S F-12 medium (GIBCO) containing 4 mM Hydrocortisone and 10% fetal calf serum. After culturing for three to four days, the cultured cells were washed with 260  $\mu$ l of Ham'S F-12 medium (GIBCO), and then added with 80  $\mu$ l of Ham'S F-12 medium containing 1 mM isobutyl-1-methyl xanthine (IBMX, SIGMA), 10% fetal calf serum and 10 mM HEPES. The resulting mixture was incubated at 37 °C for 30 min.

**[0274]** The culture mediums of the mouse antibodies, the chimeric antibodies and the humanized antibodies to be tested for neutralizing activity were previously diluted serially in the following dilution series: [10  $\mu$ g/ml, 3.3  $\mu$ g/ml, 1.1

5  $\mu\text{g/ml}$  and  $0.37 \mu\text{g/ml}$ ], [ $10 \mu\text{g/ml}$ ,  $2 \mu\text{g/ml}$ ,  $0.5 \mu\text{g/ml}$  and  $0.01 \mu\text{g/ml}$ ] and [ $10 \mu\text{g/ml}$ ,  $5 \mu\text{g/ml}$ ,  $1.25 \mu\text{g/ml}$ ,  $0.63 \mu\text{g/ml}$  and  $0.31 \mu\text{g/ml}$ ]. Each of the diluted antibody sample solutions was mixed with an equivalent amount of  $4 \text{ ng/ml}$  of PTHrP (1-34). The resulting mixed solution ( $80 \mu\text{l}$ ) was added to each well. In each well, the final concentration of each antibody became a quarter of the above-mentioned concentration of the antibody, and accordingly the concentration of PTHrP (1-34) became  $1 \text{ ng/ml}$ . After the treatment at room temperature for  $10 \text{ min.}$ , the culture supernatant was removed and the residue was washed with PBS three times. Subsequently, cAMP in the cells was extracted with  $100 \mu\text{l}$  of a  $0.3\%$  HCl- $95\%$  ethanol and then evaporated using a water jet aspirator to remove the HCl-ethanol. The residue was dissolved in  $120 \mu\text{l}$  of EIA buffer appended to cAMP EIA Kit (CAYMAN CHEMICAL'S) to extract the cAMP therefrom. The cAMP was determined using cAMP EIA Kit (CAYMAN CHEMICAL'S) in accordance with the instructions included in the kit. As a result, it was found that, among the humanized antibodies having the same levels of antigen-binding activity as that of the chimeric antibody, those antibodies having L-chain versions "q", "r", "s" and "t" (in which the 91-position tyrosine was replaced by isoleucine) exhibited the similar neutralizing activity to that of the chimeric antibody, and that antibody having a L-chain version "q" exhibited the strongest neutralizing activity.

#### 15 Sequence Listing Free Text

##### [0275]

20 SEQ ID NO: 1 Synthesized DNA  
 SEQ ID NO: 2 Synthesized DNA  
 SEQ ID NO: 3 Synthesized DNA  
 SEQ ID NO: 4 Synthesized DNA  
 SEQ ID NO: 5 Synthesized DNA  
 25 SEQ ID NO: 6 Synthesized DNA  
 SEQ ID NO: 7 Synthesized DNA  
 SEQ ID NO: 8 Synthesized DNA  
 SEQ ID NO: 9 Synthesized DNA  
 SEQ ID NO: 10 Synthesized DNA  
 SEQ ID NO: 11 Synthesized DNA  
 30 SEQ ID NO: 12 Synthesized DNA  
 SEQ ID NO: 13 Synthesized DNA  
 SEQ ID NO: 14 Synthesized DNA  
 SEQ ID NO: 15 Synthesized DNA  
 SEQ ID NO: 16 Synthesized DNA  
 35 SEQ ID NO: 17 Synthesized DNA  
 SEQ ID NO: 18 Synthesized DNA  
 SEQ ID NO: 19 Synthesized DNA  
 SEQ ID NO: 20 Synthesized DNA  
 SEQ ID NO: 21 Synthesized DNA  
 40 SEQ ID NO: 22 Synthesized DNA  
 SEQ ID NO: 23 Synthesized DNA  
 SEQ ID NO: 24 Synthesized DNA  
 SEQ ID NO: 25 Synthesized DNA  
 SEQ ID NO: 26 Synthesized DNA  
 45 SEQ ID NO: 27 Synthesized DNA  
 SEQ ID NO: 28 Synthesized DNA  
 SEQ ID NO: 29 Synthesized DNA  
 SEQ ID NO: 30 Synthesized DNA  
 SEQ ID NO: 31 Synthesized DNA  
 50 SEQ ID NO: 32 Synthesized DNA  
 SEQ ID NO: 33 Synthesized DNA  
 SEQ ID NO: 34 Synthesized DNA  
 SEQ ID NO: 35 Synthesized DNA  
 SEQ ID NO: 36 Synthesized DNA  
 55 SEQ ID NO: 37 Synthesized DNA  
 SEQ ID NO: 38 Synthesized DNA  
 SEQ ID NO: 39 Synthesized DNA  
 SEQ ID NO: 40 Synthesized DNA

SEQ ID NO: 41 Synthesized DNA  
SEQ ID NO: 42 Synthesized DNA  
SEQ ID NO: 43 Synthesized DNA  
SEQ ID NO: 44 Synthesized DNA

5

**[0276]** All publications, patents and patent applications cited herein are incorporated by reference in their entirety.

INDUSTRIAL APPLICABILITY

10

**[0277]** The present invention provides a stabilized pharmaceutical preparation of antibodies against parathyroid hormone related peptides. In addition, the present invention provides a pharmaceutical preparation for injection having a pain-easing action.

15

20

25

30

35

40

45

50

55

SEQUENCE LISTING

5

<110> CHUGAI SEIYAKU KABUSHIKI KAISHA

10

<120> Stabilized Antibody Composition and Pharmaceutical Preparation for Injection

15

<130> PH-1093-PCT

20

<150> JP 11-375203

<151> 1999-12-28

25

<160> 75

30

<170> PatentIn Ver. 2.0

<210> 1

35

<211> 20

<212> DNA

<213> Artificial Sequence

40

<220> .

45

<223> Synthetic DNA

<400> 1

50

aaatagccct tgaccaggca

20

55

<210> 2

<211> 38

<213> Artificial Sequence

$\langle 220 \rangle$

10                   <223> Synthetic DNA

<400> 2

15 ctggttcggc ccacctctga aggttccaga atcgatag

38

〈210〉 3

&lt;211&gt; 28

<212> DNA

25 <213> Artificial Sequence

 $\langle 220 \rangle$ 

30                    <223> Synthetic DNA

$\langle 400 \rangle$  3

35           ggatcccggg ccagtgkata gacagatg

28

〈210〉 4

&lt;211&gt; 29

&lt;212&gt; DNA

45 <213> Artificial Sequence

<220>

50                    <223> Synthetic DNA

<400> 4

55 ggatcccggg tcagrggaag gtggraaca

29

5           <210> 5  
            <211> 17  
            <212> DNA  
10           <213> Artificial Sequence

15       $\langle 220 \rangle$   
          $\langle 223 \rangle$  Synthetic DNA

20 <400> 5  
gttttccag tcacgac 17

25           <210> 6  
              <211> 17  
              <212> DNA  
30           <213> Artificial Sequence

35

<220>  
<223> Synthetic DNA

40 <400> 6  
caggaaacag ctatgac 17

45           <210> 7  
            <211> 31  
            <212> DNA  
50           <213> Artificial Sequence

55                      <220>



5 <223> Synthetic DNA  
 <400> 7  
 10 gtclaagcct ccacatgaa acttcgggc c 31  
 <210> 8  
 15 <211> 30  
 <212> DNA  
 <213> Artificial Sequence  
 20  
 <220>  
 <223> Synthetic DNA  
 25  
 <400> 8  
 30 tgttggatcc ctgcagagac agtgaccaga 30  
 <210> 9  
 35 <211> 36  
 <212> DNA  
 <213> Artificial Sequence  
 40  
 <220>  
 <223> Synthetic DNA  
 45  
 <400> 9  
 50 gtctgaatic aagcttcac catggggctt gggctg 36  
 <210> 10  
 55 <211> 41

5 <212> DNA  
 <213> Artificial Sequence  
  
 10 <220>  
 <223> Synthetic DNA  
  
 15 <400> 10  
 ttccccgggc ccttggtgga ggctgaggag acggtgacca g 41  
  
 20 <210> 11  
 <211> 109  
 <212> DNA  
 25 <213> Artificial Sequence  
  
 30 <220>  
 <223> Synthetic DNA  
  
 35 <400> 11  
 gtcigaattc aagcttagta cttggccagc ccaaggccaa ccccaaggic accctgttcc 60  
 cgccctcttc tgaggagctc caagccaaca aggccacact agtgtgtct 109  
 40  
 45 <210> 12  
 <211> 110  
 <212> DNA  
 <213> Artificial Sequence  
 50  
 <220>  
 <223> Synthetic DNA  
 55

5           <400> 12  
           ggtttggtag tctccactcc cgccttgacg gggctgccat ctccttcca ggccactgic 60  
           acagctcccg ggtagaagtc actgacaga cacactagtg tggccttgtt           110  
  
 10           <210> 13  
           <211> 98  
 15           <212> DNA  
           <213> Artificial Sequence  
  
 20           <220>  
           <223> Synthetic DNA  
  
 25           <400> 13  
           ggagtggaga ccaccaaac ctcacaacag agcaacaaca agtacgcggc cagcagctac 60  
 30           ctgagcctga cccccgagca gtggaagtc cacagaag           98  
  
           <210> 14  
 35           <211> 106  
           <212> DNA  
           <213> Artificial Sequence  
  
 40           <220>  
           <223> Synthetic DNA  
  
 45           <400> 14  
 50           tgttgaattc ttactatgaa catctgtag gggccactgt cttctccacg gtgtccctt 60  
           catgcgtgac ctggcagctg tagcttcgtt gggacttcca ctgctc           106  
  
 55           <210> 15

<211> 43

<212> DNA

### <213> Artificial Sequence

 $\langle 220 \rangle$ 

## <223> Synthetic DNA

<400> 15

gtctgaattc aagcttagta ctltggccagc ccaaggccaa ccc

<210> 16

<211> 20

## <212> DNA

### <213> Artificial Sequence

<220>

## <223> Synthetic DNA

<400> 16

tgttgaattc ttactatgaa

<210> 17

<211> 39

## <212> DNA

### <213> Artificial Sequence

 $\langle 220 \rangle$ 

## <223> Synthetic DNA

<400> 17  
 5 caacaagtac gcggccagca gctacctgag cctgacgcc 39  
  
 <210> 18  
 10 <211> 39  
 <212> DNA  
 15 <213> Artificial Sequence  
  
 <220>  
 20 <223> Synthetic DNA  
  
 <400> 18  
 25 gtagctgctg gccgcgtact tglgttgct ctgtttgga 39  
  
 <210> 19  
 30 <211> 46  
 <212> DNA  
 35 <213> Artificial Sequence  
  
 <220>  
 40 <223> Synthetic DNA  
  
 <400> 19  
 45 gtctgaattc aagcttagtc ctaggtcgaa ctgtggctgc accatc 46  
  
 <210> 20  
 50 <211> 34  
 <212> DNA  
 55 <213> Artificial Sequence

	<220>	
5	<223> Synthetic DNA	
	<400> 20	
10	tggtgaattc ttactaacac tctccctgt tgaa	34
15	<210> 21	
	<211> 35	
	<212> DNA	
20	<213> Artificial Sequence	
25	<220>	
	<223> Synthetic DNA	
30	<400> 21	
	gtctaagctt ccaccaiggc ctggactcct ctctt	35
35	<210> 22	
	<211> 48	
40	<212> DNA	
	<213> Artificial Sequence	
45	<220>	
	<223> Synthetic DNA	
50	<400> 22	
	tggtgaattc agatctaact acttacctag gacagtgacc ttggcacc	48
55		

<210> 23

<211> 128

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA

<400> 23

gtctaagctt ccaccaagg gtttgggctg agctgggtt: tcttcgttgc tcttttaaga 60  
gggtgtccagt gtcagggtgca gctgggtggag tctgggggag gcgtgggtcca gccctgggagg 120  
tccctgag 128

<210> 24

<211> 125

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic DNA

<400> 24

accattagta gttgtggttag ttacacctac tatccagaca gtgtgaaggg gcgattcacc 60  
atctccagag acaattccaa gaacacgctg tatctgcaaa tgaacagcct gagagctgag 120  
gacac 125

<210> 25

<211> 132

<212> DNA

<213> Artificial Sequence

5

<220>

<223> Synthetic DNA

10

<400> 25

15

ctaccaccac tactaatggt tgcacccac tccagcccct tgcctggagc ctggcggacc 60  
caagacatgc catagctact gaaggigaat ccagaggctg cacaggagag tctcaggac 120  
ctcccaggct gg 132

20

<210> 26

<211> 110

25

<212> DNA

<213> Artificial Sequence

30

<220>

<223> Synthetic DNA

35

<400> 26

40

igtggatcc ctgaggagac ggtgaccagg gtccctggc cccagtaagc aaagtaagtc 60  
atagtagtct gtcctgcaca gtaatacaca gccggtcct cagctctcag 110

45

<210> 27

<211> 30

<212> DNA

50

<213> Artificial Sequence

<220>

55

<223> Synthetic DNA



5                   <400> 27  
                  gtctaaagctt ccaccatggg gtttgggctg                   30

10	<210> 28
	<211> 30
	<212> DNA
15	<213> Artificial Sequence

20       $\langle 220 \rangle$   
           $\langle 223 \rangle$  Synthetic DNA

25 <400> 28  
tgttggatcc ctgaggagac ggtgaccagg 30

30           <210> 29  
            <211> 133  
35           <212> DNA  
            <213> Artificial Sequence

40           <220>  
              <223> Synthetic DNA

45 <400> 29  
 acaaagcctc caccatggcc tggactccctc tcttctctct ctttgttctt cattgctcag 60  
 50 gttctttctc ccagcttgtg ctgactcaat cgccctctgc ctctgcctcc ctgggagcct 120  
 cggatcaagct cac 133

55  $\langle 210 \rangle$  30

&lt;211&gt; 118

5                     $\langle 212 \rangle$  DNA

### <213> Artificial Sequence

10.  $\langle 220 \rangle$

## <223> Synthetic DNA

15

$\langle 400 \rangle$  30

20 agcaagatgg aagccacagc acaggtgatg ggattcciga tgccttctca ggctccagct 60  
ctggggctga gcgctacctc accatctcca gccctcagtc tgaggatgag gctgacta 118

&lt;210&gt; 31

&lt;211&gt; 128

## <212> DNA

30 <213> Artificial Sequence

 $\langle 220 \rangle$ 

35           &lt;223&gt; Synthetic DNA

<400> 31

40      ctgtggcttc cactttgctt aagtttcaic aagtaccgag ggcccttctc tggctgctgc 60  
tgatgccatt caatgggtga cgtactgtgc tgactactca aggtgcaggt gagcttgacc 120  
gaggctcc 128

&lt;210&gt; 32

&lt;211&gt; 114

## <212> DNA

### <213> Artificial Sequence

5

10

15

20

25

30

35

40

45

50

55

5 <210> 35  
 <211> 75  
 <212> DNA  
 <213> Artificial Sequence  
 10  
 <220>  
 <223> Synthetic DNA  
 15  
 <400> 35  
 20 ctltggatccg ggctgacctt ggacggctcag tttaggtcct ccgccgaaca cgtacacaaa 60  
 ttgttcctta attgt 75  
 25 <210> 36  
 <211> 43  
 <212> DNA  
 30 <213> Artificial Sequence  
 <220>  
 35 <223> Synthetic DNA  
 <400> 36  
 40 aaaggatcct taagatccat caagtaccga gggggcttct ctg 43  
 45 <210> 37  
 <211> 46  
 <212> DNA  
 50 <213> Artificial Sequence  
 55 <220>

5 <223> Synthetic DNA

<400> 37

10 acaaagctta ggcctacctc accatctcca gccctccagcc tgagga 46

<210> 38

15 <211> 111

<212> DNA

<213> Artificial Sequence

20

<220>

25 <223> Synthetic DNA

<400> 38

30 cttggatccg ggctgacctt ggacggctcag ttgggtccct ccgccgaaca cgtacacaaa 60

ttgttcctta attgtatcac ccacaccaca gatatagtcg gccctcctc c 111

35 <210> 39

<211> 42

<212> DNA

40 <213> Artificial Sequence

<220>

45 <223> Synthetic DNA

<400> 39

50 cttctctggc tgctgctgat accattcaat ggtglacgta ct 42

55 <210> 40

5

10

15

	<400> 42	
5	cacgaattca ctatcgattc tggaaaccttc agagg	35
10	<210> 43	
	<211> 18	
	<212> DNA	
15	<213> Artificial Sequence	
20	<220>	
	<223> Synthetic DNA	
25	<400> 43	
	ggccttgagc tcctcaga	18
30	<210> 44	
	<211> 20	
	<212> DNA	
35	<213> Artificial Sequence	
40	<220>	
	<223> Synthetic DNA	
45	<400> 44	
	gacagtgggtt caaagttttt	20
50	<210> 45	
	<211> 118	
	<212> PRT	
55	<213> Mus musculus	

5

&lt;400&gt; 45

10

Gln Leu Val Leu Thr Gln Ser Ser Ser Ala Ser Phe Ser Leu Gly Ala

1 5 10 15

Ser Ala Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr

20 25 30

15

Ile Glu Trp Tyr Gln Gln Gln Pro Leu Lys Pro Pro Lys Tyr Val Met

35 40 45

20

Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp

50 55 60

25

Arg Phe Ser Gly Ser Ser Ser Gly Ala Asp Arg Tyr Leu Ser Ile Ser

65 70 75 80

Asn Ile Gln Pro Glu Asp Glu Ala Met Tyr Ile Cys Gly Val Gly Asp

85 90 95

30

Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Val

100 105 110

Thr Val Leu Gly Gln Pro

35

115

40

&lt;210&gt; 46

&lt;211&gt; 118

&lt;212&gt; PRT

45

&lt;213&gt; Mus musculus

50

&lt;400&gt; 46

Glu Val Gln Leu Val Glu Ser Gly Gly Asp Leu Val Lys Pro Gly Gly

1 5 10 15

Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr

55

20 25 30



EP 1 254 666 A1

Gly Met Ser Trp Ile Arg Gln Thr Pro Asp Lys Arg Leu Glu Trp Val  
 35 40 45  
 5 Ala Thr Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Pro Asp Ser Val  
 50 55 60  
 10 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Met Phe Tyr Cys  
 15 85 90 95  
 Ala Arg Gln Thr Thr Met Thr Tyr Phe Ala Tyr Trp Gly Gln Gly Thr  
 100 105 110  
 20 Leu Val Thr Val Ser Ala  
 115  
 25  
 <210> 47  
 <211> 116  
 30 <212> PRT  
 <213> Homo sapiens  
 35  
 <400> 47  
 Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser Leu Gly Ala  
 40 1 5 10 15  
 Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr  
 20 25 30  
 45 Ile Glu Trp His Gln Gln Gln Pro Glu Lys Gly Pro Arg Tyr Leu Met  
 35 40 45  
 Lys Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp  
 50 55 60  
 Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser  
 55 65 70 75 80

5 Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Val Gly Asp  
85 90 95  
Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Leu  
10 100 105 110  
Thr Val Leu Gly  
115

15

<210> 48  
20 <211> 118  
<212> PRT  
<213> Homo sapiens  
25

<400> 48  
30 Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser Leu Gly Ala  
1 5 10 15  
Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr  
35 20 25 30  
Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Lys Tyr Leu Met  
35 40 45  
40 Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp  
50 55 60  
Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser  
45 65 70 75 80  
Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Val Gly Asp  
50 85 90 95  
Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Leu  
100 105 110  
55 Thr Val Leu Gly Gln Pro

115

5

&lt;210&gt; 49

&lt;211&gt; 118

10

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

15

&lt;400&gt; 49

Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser Leu Gly Ala

20

1 5 10 15

Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr

20 25 30

25

Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Lys Tyr Val Met

35 40 45

30

Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp

50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser

35

65 70 75 80

Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Val Gly Asp

85 90 95

40

Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Leu

100 105 110

Thr Val Leu Gly Gln Pro

45

115

50

&lt;210&gt; 50

&lt;211&gt; 118

&lt;212&gt; PRT

55

&lt;213&gt; Homo sapiens

5

&lt;400&gt; 50

10

Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser Leu Gly Ala

1 5 10 15

Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr

20 25 30

15

Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Arg Tyr Leu Met

35 40 45

20

Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp

50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser

65 70 75 80

25

Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Val Gly Asp

85 90 95

30

Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Leu

100 105 110

Thr Val Leu Gly Gln Pro

35

115

40

&lt;210&gt; 51

&lt;211&gt; 118

&lt;212&gt; PRT

45

&lt;213&gt; Homo sapiens

50

&lt;400&gt; 51

Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser Leu Gly Ala

1 5 10 15

Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr

55

20 25 30

EP 1 254 666 A1

Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Arg Tyr Val Met  
 5                   35                   40                   45  
 Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp  
                   50                   55                   60  
 10 Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser  
      65                   70                   75                   80  
 Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Val Gly Asp  
 15                   85                   90                   95  
 Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Leu  
 20                   100                   105                   110  
 Thr Val Leu Gly Gln Pro  
                   115  
 25  
 <210> 52  
 <211> 118  
 30 <212> PRT  
 <213> Homo sapiens  
 35  
 <400> 52  
 Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser Leu Gly Ala  
 40       1                   5                   10                   15  
 Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr  
                   20                   25                   30  
 45 Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Lys Tyr Leu Met  
                   35                   40                   45  
 50 Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp  
                   50                   55                   60  
 Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser  
 55       65                   70                   75                   80

EP 1 254 666 A1

Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly Val Gly Asp  
 5                               85                               90                               95  
 Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Leu  
                              100                               105                               110  
 10 Thr Val Leu Gly Gln Pro  
                              115  
 15  
 <210> 53  
 <211> 118  
 20 <212> PRT  
 <213> Homo sapiens  
 25  
 <400> 53  
 Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser Leu Gly Ala  
      1                               5                               10                               15  
 30 Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr  
                              20                               25                               30  
 Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Arg Tyr Leu Met  
 35                               35                               40                               45  
 Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp  
                              50                               55                               60  
 40 Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser  
      65                               70                               75                               80  
 45 Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly Val Gly Asp  
                              85                               90                               95  
 Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Leu  
 50                               100                               105                               110  
 Thr Val Leu Gly Gln Pro  
 55                               115

5 <210> 54  
 <211> 118  
 <212> PRT  
 10 <213> Homo sapiens

<400> 54  
 15 Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser Leu Gly Ala  
 1 5 10 15  
 20 Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr  
 20 25 30  
 Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Lys Tyr Val Met  
 25 35 40 45  
 Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp  
 50 55 60  
 30 Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser  
 65 70 75 80  
 35 Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly Val Gly Asp  
 85 90 95  
 Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Leu  
 40 100 105 110  
 Thr Val Leu Gly Gln Pro  
 115

45  
 <210> 55  
 <211> 118  
 50 <212> PRT  
 <213> Homo sapiens  
 55

&lt;400&gt; 55

5 Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser Leu Gly Ala  
     1                    5                    10                    15  
 Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser Thr Tyr Thr  
 10                    20                    25                    30  
 Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Arg Tyr Val Met  
                     35                    40                    45  
 15 Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp  
                     50                    55                    60  
 20 Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser  
                     65                    70                    75                    80  
 Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly Val Gly Asp  
 25                    85                    90                    95  
 Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly Thr Lys Leu  
                     100                    105                    110  
 30 Thr Val Leu Gly Gln Pro  
                     115

35

&lt;210&gt; 56

&lt;211&gt; 118

40

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

45

&lt;400&gt; 56

50 Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro Gly Arg  
     1                    5                    10                    15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr  
                     20                    25                    30  
 55 Gly Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val



EP 1 254 666 A1

35 40 45  
 5 Ala Thr Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Pro Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 10 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 15 Ala Arg Gln Thr Thr Met Thr Tyr Phe Ala Tyr Trp Gly Gln Gly Thr  
 100 105 110  
 20 Leu Val Thr Val Ser Ser  
 115  
 25 <210> 57  
 <211> 411  
 <212> DNA  
 30 <213> Mus musculus  
 35 <220>  
 <221> CDS  
 <222> (1).. (411)  
 40 <220>  
 <221> mat\_peptide  
 45 <222> (58).. (411)  
 50 <400> 57  
 'aig aac ttc ggg ctc agc ttg att ttc ctt gcc ctc att tta aaa ggt 48  
 Met Asn Phe Gly Leu Ser Leu Ile Phe Leu Ala Leu Ile Leu Lys Gly  
 55 -15 -10 -5

EP 1 254 666 A1

5 gtc cag tgt gag gtg caa ctg gtc gag tct ggg gga gac tta gtg aag 96  
 Val Gln Cys Glu Val Gln Leu Val Glu Ser Gly Gly Asp Leu Val Lys  
 -1 1 5 10  
 10 cct gga ggg tcc ctg aaa ctg tcc tgt gca gcc tct gga ttc act ttc 144  
 Pro Gly Gly Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe  
 15 20 25  
 15 agt agc tat ggc atg tct tgg att cgc cag act cca gac aag agg ctg 192  
 Ser Ser Tyr Gly Met Ser Trp Ile Arg Gln Thr Pro Asp Lys Arg Leu  
 30 35 40 45  
 20 gag tgg gtc gca acc att agt agt ggt ggt agt tac acc tac tat cca 240  
 Glu Trp Val Ala Thr Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Pro  
 50 55 60  
 25 gac agt gtc aag ggg cga ttc acc atc tcc aga gac aat gcc aag aac 288  
 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn  
 65 70 75  
 30 acc cta tac ctg caa atg agc agt ctg aag tct gag gac aca gcc atg 336  
 Thr Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Met  
 80 85 90  
 35 ttt tac tgt gca aga cag act act atg act tac ttt gct tac tgg ggc 384  
 Phe Tyr Cys Ala Arg Gln Thr Thr Met Thr Tyr Phe Ala Tyr Trp Gly  
 95 100 105  
 40 caa ggg act ctg gtc act gtc tct gca 411  
 Gln Gly Thr Leu Val Thr Val Ser Ala  
 45 110 115  
 50 <210> 58  
 <211> 411  
 <212> DNA  
 55 <213> Homo sapiens

5 <220>  
 <221> CDS  
 <222> (1).. (411)

10 <220>  
 <221> mat\_peptide  
 15 <222> (58).. (411)

20 <400> 58  
 atg ggg ttt ggg ctg agc tgg gtt ttc ctc gtt gct ctt tta aga ggt 48  
 Met Gly Phe Gly Leu Ser Trp Val Phe Leu Val Ala Leu Leu Arg Gly  
 25 -15 -10 -5  
 gtc cag tgt cag gtg cag ctg gtg gag tct ggg gga ggc gtg gtc cag 96  
 Val Gln Cys Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln  
 30 -1 1 5 10  
 cct ggg agg tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc 144  
 Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe  
 35 15 20 25  
 agt agc tat ggc atg tct tgg gtc cgc cag gct cca ggc aag ggg ctg 192  
 Ser Ser Tyr Gly Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu  
 40 30 35 40 45  
 gag tgg gtg gca acc att agt agt ggt ggt agt tac acc tac tat cca 240  
 Glu Trp Val Ala Thr Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Pro  
 45 50 55 60  
 gac agt gtg aag ggg cga ttc acc atc tcc aga gac aat tcc aag aac 288  
 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn  
 50 65 70 75  
 55 acg ctg tat ctg caa atg aac agc ctg aga gct gag gac acg gct gtg 336

5 Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val  
 80 85 90  
 tat tac tgt gcg aga cag act act atg act tac ttt gct tac tgg ggc 384  
 10 Tyr Tyr Cys Ala Arg Gln Thr Thr Met Thr Tyr Phe Ala Tyr Trp Gly  
 95 100 105  
 cag gga acc ctg gtc acc gtc tcc tca 411  
 15 Gln Gly Thr Leu Val Thr Val Ser Ser  
 110 115  
 20 <210> 59  
 <211> 11  
 25 <212> PRT  
 <213> Homo sapiens  
 30 <400> 59  
 Lys Ala Ser Gln Asp Val Asn Thr Ala Val Ala  
 1 5 10  
 35 <210> 60  
 <211> 7  
 40 <212> PRT  
 <213> Homo sapiens  
 45 <400> 60  
 Ser Ala Ser Asn Arg Tyr Thr  
 50 1 5  
 55 <210> 61  
 <211> 9

5           <212> PRT  
           <213> Homo sapiens  
  
 10           <400> 61  
           Gln Gln His Tyr Ser Thr Pro Phe Thr  
           1                   5  
  
 15           <210> 62  
           <211> 5  
 20           <212> PRT  
           <213> Homo sapiens  
  
 25           <400> 62  
           Pro Tyr Trp Met Gln  
 30           1                   5  
  
           <210> 63  
 35           <211> 16  
           <212> PRT  
           <213> Homo sapiens  
  
 40           <400> 63  
 45           Ser Ile Phe Gly Asp Gly Asp Thr Arg Tyr Ser Gln Lys Phe Lys Gly  
           1                   5                   10                   15  
  
 50           <210> 64  
           <211> 11  
           <212> PRT  
 55           <213> Homo sapiens

5 <400> 64  
 Gly Leu Arg Arg Gly Gly Tyr Tyr Phe Asp Tyr  
 1 5 10  
 10  
 <210> 65  
 15 <211> 411  
 <212> DNA  
 <213> Mus musculus  
 20  
 <220>  
 <221> CDS  
 25 <222> (1).. (411)  
 <220>  
 30 <221> mat\_peptide  
 <222> (58).. (411)  
 35  
 <400> 65  
 atg gcc tgg act cct ctc ttc ttc ttt gtt ctt cat tgc tca ggt 48  
 40 Met Ala Trp Thr Pro Leu Phe Phe Phe Phe Val Leu His Cys Ser Gly  
 -15 -10 -5  
 tct ttc tcc caa ctt gtg ctc act cag tca tct tca gcc tct ttc tcc 96  
 45 Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Ser Ser Ala Ser Phe Ser  
 -1 1 5 10  
 50 ctg gga gcc tca gca aaa ctc acg tgc acc ttg agt agi cag cac agt 144  
 Leu Gly Ala Ser Ala Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser  
 15 20 25  
 55 acg tac acc att gaa tgg tat cag caa cag cca ctc aag cct cct aag 192

Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Leu Lys Pro Pro Lys  
 5           30                   35                   40                   45  
 tat gtg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg 240  
 Tyr Val Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly  
 10                   50                   55                   60  
 att cct gat cgc ttc tct gga tcc agc tct ggt gct gat cgc tac ctt 288  
 Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Asp Arg Tyr Leu  
 15                   65                   70                   75  
 agc att tcc aac atc cag cca gaa gat gaa gca atg tac atc tgt ggt 336  
 Ser Ile Ser Asn Ile Gln Pro Glu Asp Glu Ala Met Tyr Ile Cys Gly  
 20                   80                   85                   90  
 gtg ggt gat aca att aag gaa caa ttt gtg tat gtt ttc ggc ggt ggg 384  
 25 Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly  
 95                   100                   105  
 acc aag gtc act gtc cta ggt cag ccc 411  
 30 Thr Lys Val Thr Val Leu Gly Gln Pro  
 110                   115  
 35  
 <210> 66  
 <211> 411  
 40 <212> DNA  
 <213> Homo sapiens  
 45  
 <220>  
 <221> CDS  
 50 <222> (1).. (411)  
 <220>  
 55 <221> mat\_peptide

&lt;222&gt; (58).. (411)

5

&lt;400&gt; 66

10

atg gcc tgg act cct ctc ttc ttc ttc ttt gtt ctt cat tgc tca ggt 48

Met Ala Trp Thr Pro Leu Phe Phe Phe Phe Val Leu His Cys Ser Gly

-15

-10

-5

15

tct ttc tcc cag ctt gtg ctg act caa tgc ccc tct gcc tct gcc tcc 96

Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser

-1 1

5

10

20

ctg gga gcc tgc gtc aag ctc acc tgc acc ttg agt agt cag cac agt 144

Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser

15

20

25

25

acg tac acc att gaa tgg cat cag cag cag cca gag aag ggc cct cgg 192

Thr Tyr Thr Ile Glu Trp His Gln Gln Gln Pro Glu Lys Gly Pro Arg

30

35

40

45

30

tac ttg atg aaa ctt aag caa gat gga agc cac agc aca ggt gat ggg 240

Tyr Leu Met Lys Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly

50

55

60

35

att cct gat cgc ttc tca ggc tcc agc tct ggg gct gag cgc tac ctc 288

Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu

65

70

75

40

acc atc tcc agc ctc cag tct gag gat gag gct gac tat tac tgt ggt 336

Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly

80

85

90

45

gtg ggt gat aca att aag gaa caa ttt gtg tac glg ttc ggc gga ggg 384

Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly

95

100

105

50

acc aaa ctg acc gtc cta ggt cag ccc 411

55

Thr Lys Leu Thr Val Leu Gly Gln Pro



110 115

5  
 <210> 67  
 <211> 411  
 10  
 <212> DNA  
 <213> Homo sapiens

15  
 <220>  
 <221> CDS  
 20  
 <222> (1).. (411)

<220>  
 25  
 <221> mat\_peptide  
 <222> (58).. (411)

30  
 <400> 67  
 atg gcc tgg act cct ctc ttc ttc ttc ttt gtt ctt cat tgc tca ggt 48  
 35 Met Ala Trp Thr Pro Leu Phe Phe Phe Phe Val Leu His Cys Ser Gly  
 -15 -10 -5  
 tct ttc tcc cag ctt gtc ctg act caa tgc ccc tct gcc tct gcc tcc 96  
 40 Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser  
 -1 1 5 10  
 ctg gga gcc tgc gtc aag ctc acc tgc acc ttg agt agt cag cac agt 144  
 45 Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser  
 15 20 25  
 acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct aag 192  
 50 Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Lys  
 30 35 40 45  
 55 tac ctg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg 240

Tyr Leu Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly  
 5                                      50                                      55                                      60  
 att cct gat cgc ttc tca ggc tcc agc tct ggg gct gag cgc tac ctc    288  
 10    Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu  
                                     65                                      70                                      75  
 acc atc tcc agc ctc cag tct gag gat gag gct gac tat tac tgt ggt    336  
 15    Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly  
                                     80                                      85                                      90  
 gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggg    384  
 20    Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly  
                                     95                                      100                                      105  
 acc aaa ctg acc gtc cta ggc cag ccc                                      411  
 25    Thr Lys Leu Thr Val Leu Gly Gln Pro  
                                     110                                      115  
 30  
     <210> 68  
     <211> 411  
 35    <212> DNA  
     <213> Homo sapiens  
 40  
     <220>  
     <221> CDS  
     <222> (1).. (411)  
 45  
     <220>  
     <221> mat\_peptide  
     <222> (58).. (411)  
 50  
 55    <400> 68

atg gcc tgg act cct ctc ttc ttc ttt gtt ctt cat tgc tca ggt 48  
 Met Ala Trp Thr Pro leu Phe Phe Phe Phe Val Leu His Cys Ser Gly  
 5                   -15                   -10                   -5  
 tct ttc tcc cag ctt gtg ctg act caa tgc ccc tct gcc tct gcc tcc 96  
 Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser  
 10                   -1    1                   5                   10  
 ctg gga gcc tgc gtc aag ctc acc tgc acc ttg agt agt cag cac agt 144  
 Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser  
 15                   15                   20                   25  
 acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct aag 192  
 Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Lys  
 20                   30                   35                   40                   45  
 tac gtg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg 240  
 Tyr Val Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly  
 25                   50                   55                   60  
 att cct gat cgc ttc tca ggc tcc agc tct ggg gct gag cgc tac ctc 288  
 Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu  
 30                   65                   70                   75  
 acc atc tcc agc ctc cag tct gag gat gag gct gac tat tac tgt ggt 336  
 Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly  
 35                   80                   85                   90  
 gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggg 384  
 Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly  
 40                   95                   100                   105  
 acc aaa ctg acc gtc cta ggc cag ccc 411  
 Thr Lys Leu Thr Val Leu Gly Gln Pro  
 45                   110                   115  
 50  
 55 <210> 69

<211> 411  
 5 <212> DNA  
 <213> Homo sapiens  
 10 <220>  
 <221> CDS  
 15 <222> (1).. (411)  
 <220>  
 20 <221> mat\_peptide  
 <222> (58).. (411)  
 25 <400> 69  
 atg gcc tgg act cct ctc ttc ttc ttc ttt gtt ctt cat tgc tca ggt 48  
 Met Ala Trp Thr Pro Leu Phe Phe Phe Phe Val Leu His Cys Ser Gly  
 30 -15 -10 -5  
 tct ttc tcc cag ctt gtg ctg act caa tgc ccc tct gcc tct gcc tcc 96  
 Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser  
 35 -1 1 5 10  
 ctg gga gcc tgc gtc aag ctc acc tgc acc ttg agt agt cag cac agt 144  
 Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser  
 40 15 20 25  
 acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct agg 192  
 Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Arg  
 45 30 35 40 45  
 tac ctg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg 240  
 Tyr Leu Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly  
 50 50 55 60  
 55 att cct gat cgc ttc tca ggc tcc agc tct ggg gct gag cgc tac ctc 288

Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu  
 5                      65                      70                      75  
 acc atc tcc agc ctc cag tct gag gat gag gct gac tat tac tgt ggt 336  
 Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly  
 10                      80                      85                      90  
 gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggg 384  
 Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly  
 15                      95                      100                      105  
 acc aaa ctg acc gtc cta ggc cag ccc 411  
 20 Thr Lys Leu Thr Val Leu Gly Gln Pro  
 110                      115  
 25 <210> 70  
 <211> 411  
 <212> DNA  
 30 <213> Homo sapiens  
 35 <220>  
 <221> CDS  
 <222> (1).. (411)  
 40 <220>  
 <221> mat\_peptide  
 45 <222> (58).. (411)  
 50 <400> 70  
 atg gcc tgg act cct ctc ttc ttc ttc ttt gtt ctt cat tgc tca ggt 48  
 Met Ala Trp Thr Pro Leu Phe Phe Phe Phe Val Leu His Cys Ser Gly  
 55                      -15                      -10                      -5

EP 1 254 666 A1

5           tct ttc tcc cag ctt glg ctg act caa tgg ccc tct gcc tct gcc tcc   96  
       Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser  
               -1   1                       5                       10  
 10       ctg gga gcc tgg gtc aag ctg acc tgc acc ttg agt agt cag cac agt   144  
       Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser  
               15                       20                       25  
 15       acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct agg   192  
       Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Arg  
               30                       35                       40                       45  
 20       tac glg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg   240  
       Tyr Val Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly  
                               50                       55                       60  
 25       att cct gat cgc ttc tca ggc tcc agc tct ggg gct gag cgc tac ctg   288  
       Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu  
                               65                       70                       75  
 30       acc atc tcc agc ctg cag tct gag gat gag gct gac tat tac tgt ggt   336  
       Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly  
                               80                       85                       90  
 35       gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggg   384  
       Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly  
                               95                       100                       105  
 40       acc aaa ctg acc gtc ctg ggc cag ccc                       411  
       Thr Lys Leu Thr Val Leu Gly Gln Pro  
 45       110                       115  
  
 50       <210> 71  
       <211> 411  
       <212> DNA  
 55       <213> Homo sapiens

<220>  
 5 <221> CDS  
 <222> (1).. (411)

10 <220>  
 <221> mat\_peptide  
 15 <222> (58).. (411)

<400> 71

20 atg gcc tgg act cct ctc ttc ttc ttc ttt gtt ctt cat tgc tca ggt 48  
 Met Ala Trp Thr Pro Leu Phe Phe Phe Phe Val Leu His Cys Ser Gly  
 -15 -10 -5

25 tct ttc tcc cag ctt gtg ctg act caa tgc ccc tct gcc tct gcc tcc 96  
 Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser  
 -1 1 5 10

30 ctg gga gcc tgc gtc aag ctc acc tgc acc ttg agt agt cag cac agt 144  
 Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser  
 15 20 25

35 acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct aag 192  
 Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Lys  
 30 35 40 45

40 tac ctg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg 240  
 Tyr Leu Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly  
 50 55 60

45 att cct gat cgc ttc tca ggc tcc agc tct ggg gct gag cgc tac ctc 288  
 Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu  
 65 70 75

50 acc atc tcc agc ctc cag tct gag gat gag gct gac tat atc tgt ggt 336

Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly  
 5                      80                      85                      90  
 gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggg 384  
 Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly  
 10                      95                      100                      105  
 acc aaa ctg acc gtc cta ggc cag ccc 411  
 15 Thr Lys Leu Thr Val Leu Gly Gln Pro  
 110                      115  
 20 <210> 72  
 <211> 411  
 <212> DNA  
 25 <213> Homo sapiens  
 30 <220>  
 <221> CDS  
 35 <222> (1).. (411)  
 40 <220>  
 <221> mat\_peptide  
 <222> (58).. (411)  
 45 <400> 72  
 atg gcc lgg act cct ctc ttc ttc ttc ttt gtt ctt cat tgc tca ggt 48  
 Met Ala Trp Thr Pro Leu Phe Phe Phe Phe Val Leu His Cys Ser Gly  
 50                      -15                      -10                      -5  
 tct ttc tcc cag ctt gtg ctg act caa tgg ccc tct gcc tct gcc tcc 96  
 Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser  
 55                      -1                      1                      5                      10



**EP 1 254 666 A1**

	ctg gga gcc tcg gtc aag ctc acc tgc acc ttg agt agt cag cac agt	144
5	Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser	
	15 20 25	
	acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct agg	192
10	Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Arg	
	30 35 40 45	
	tac ctg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg	240
15	Tyr Leu Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly	
	50 55 60	
	att cct gat cgc ttc tca ggc tcc agc tct ggg gct gag cgc tac ctc	288
20	Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu	
	65 70 75	
	acc atc tcc agc ctc cag tct gag gat gag gct gac tat atc tgt ggt	336
25	Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly	
	80 85 90	
	gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggg	384
30	Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly	
	95 100 105	
	acc aaa ctg acc gtc cta ggc cag ccc	411
35	Thr Lys Leu Thr Val Leu Gly Gln Pro	
	110 115	
40		

〈210〉 73

45  $\langle 211 \rangle$  411

## <212> DNA

<213> Homo sapiens

 $\langle 220 \rangle$ 

55 &lt;221&gt; CDS

<222> (1).. (411)

5

<220>

<221> mal\_peptide

10

<222> (58).. (411)

<400> 73

15

atg gcc tgg act cct ctc ttc ttc ttc ttt gtt ctt cat tgc tca ggt 48  
Met Ala Trp Thr Pro Leu Phe Phe Phe Phe Val Leu His Cys Ser Gly

20

-15 -10 -5  
tct ttc tcc cag ctt gtg ctg act caa tgc ccc tct gcc tct gcc tcc 96  
Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser

25

-1 1 5 10  
ctg gga gcc tgc gtc aag ctc acc tgc acc ttg agt agt cag cac agt 144  
Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser

30

15 20 25  
acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct aag 192  
Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Lys

35

30 35 40 45  
tac gtg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg 240  
Tyr Val Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly

40

50 55 60  
att cct gat cgc ttc tca ggc tcc agc tct ggg gct gag cgc tac ctc 288  
Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu

45

65 70 75  
acc atc tcc agc ctc cag tct gag gat gag gct gac tat atc tgt ggt 336  
Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly

50

80 85 90  
gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggg 384

55

Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly  
 5           95                   100                   105  
 acc aaa ctg acc gtc cta ggc cag ccc                   411  
 Thr Lys Leu Thr Val Leu Gly Gln Pro  
 10       110                   115  
  
 <210> 74  
 15       <211> 411  
       <212> DNA  
 20       <213> Homo sapiens  
  
       <220>  
 25       <221> CDS  
       <222> (1).. (411)  
  
 30       <220>  
       <221> mat\_peptide  
 35       <222> (58).. (411)  
  
       <400> 74  
 40       atg gcc tgg act cct ctc ttc ttc ttc ttt gtt ctt cat tgc tca ggt   48  
       Met Ala Trp Thr Pro Leu Phe Phe Phe Phe Val Leu His Cys Ser Gly  
                      -15                   -10                   -5  
 45       tct ttc tcc cag ctt gtg ctg act caa tgc ccc tct gcc tct gcc tcc   96  
       Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser  
                      -1    1                   5                   10  
 50       ctg gga gcc tgc gtc aag ctc acc tgc acc ttg agt agt cag cac agt   144  
       Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser  
 55       15                   20                   25

acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct agg 192  
 5 Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Arg  
 30 35 40 45  
 tac gtc atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg 240  
 10 Tyr Val Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly  
 50 55 60  
 att cct gat cgc ttc tca ggc tcc agc tct ggg gct gag cgc tac ctc 288  
 15 Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu  
 65 70 75  
 acc atc tcc agc ctc cag tct gag gat gag gct gac tat atc tgt ggt 336  
 20 Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly  
 80 85 90  
 gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggg 384  
 25 Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly  
 95 100 105  
 30 acc aaa ctg acc gtc cta ggc cag ccc 411  
 Thr Lys Leu Thr Val Leu Gly Gln Pro  
 110 115  
 35  
 <210> 75  
 40 <211> 34  
 <212> PRT  
 <213> Homo sapiens  
 45  
 <400> 75  
 50 Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly Lys Ser Ile Gln  
 1 5 10 15  
 Asp Leu Arg Arg Arg Phe Phe Leu His His Leu Ile Ala Glu Ile His  
 55 20 25 30

Thr Ala

5

## Claims

- 10 1. A stabilized pharmaceutical preparation of an antibody to parathyroid hormone related peptide, wherein the antibody is dissolved in a buffer solution containing at least one buffer selected from the group consisting of acetic acid, citric acid, phosphoric acid, and a salt thereof and is in the form of a solution of pH 5 to 8.
- 15 2. The stabilized pharmaceutical preparation according to claim 1, having a total concentration of the buffer of 0.1 to 100 mmol/L.
3. The stabilized pharmaceutical preparation according to claim 1, having a total concentration of the buffer of 5 to 50 mmol/L.
- 20 4. The stabilized pharmaceutical preparation according to any one of claims 1 to 3, wherein the antibody to parathyroid hormone related peptide is monoclonal.
5. The stabilized pharmaceutical preparation according to claim 4, wherein the antibody to parathyroid hormone related peptide is a human antibody, a humanized antibody, or a chimeric antibody.
- 25 6. A stabilized antibody solution composition to parathyroid hormone related peptide, wherein the antibody is dissolved in a buffer solution containing at least one buffer selected from the group consisting of acetic acid, citric acid, phosphoric acid, and a salt thereof and is in the form of a solution of pH 5 to 8.
- 30 7. The antibody solution composition according to claim 6, wherein the antibody solution composition is a solution composition for bulk.
8. The antibody solution composition according to claim 7, substantially free of a stabilizer other than a buffer or an isotonicizing agent.
- 35 9. A pharmaceutical preparation for injection wherein an antibody to parathyroid hormone related peptide is dissolved in a buffer solution containing a buffer consisting of acetic acid and/or a salt thereof.
- 40 10. The pharmaceutical preparation for injection according to claim 9, wherein the preparation is in the form of a solution of pH 5 to 8.
11. The pharmaceutical preparation for injection according to claim 9, wherein the preparation has a total concentration of the buffer of 0.1 to 100 mmol/L.
- 45 12. The pharmaceutical preparation for injection according to claim 9, wherein the preparation has a total concentration of the buffer of 5 to 50 mmol/L.
13. The pharmaceutical preparation for injection according to any one of claims 9 to 12, wherein the antibody to parathyroid hormone related peptide is monoclonal.
- 50 14. The pharmaceutical preparation for injection according to claim 13, wherein the antibody is a human antibody, a humanized antibody, or a chimeric antibody.

55

FIG.1

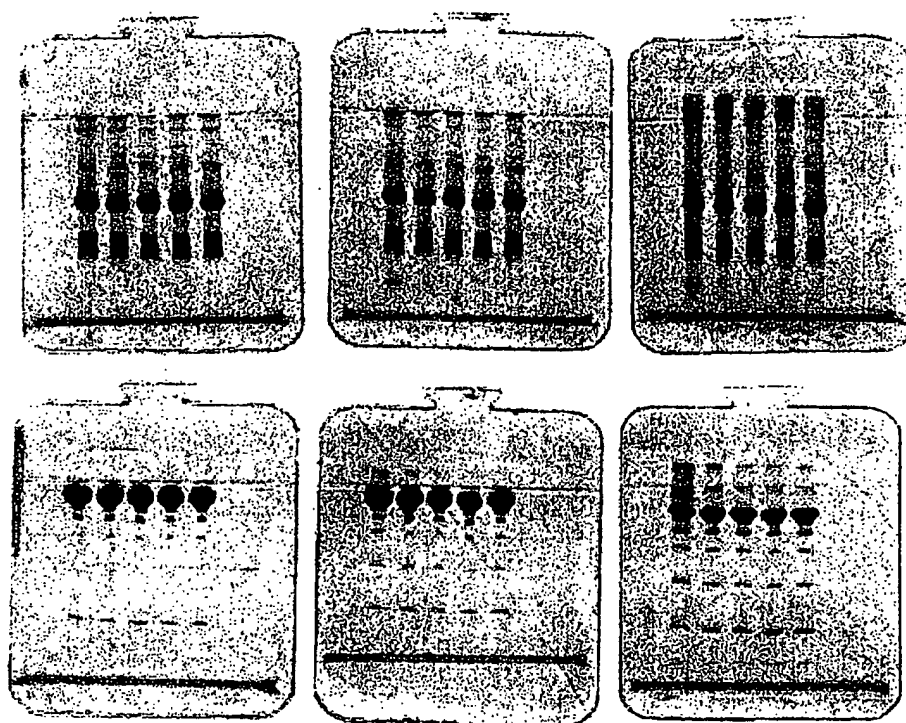
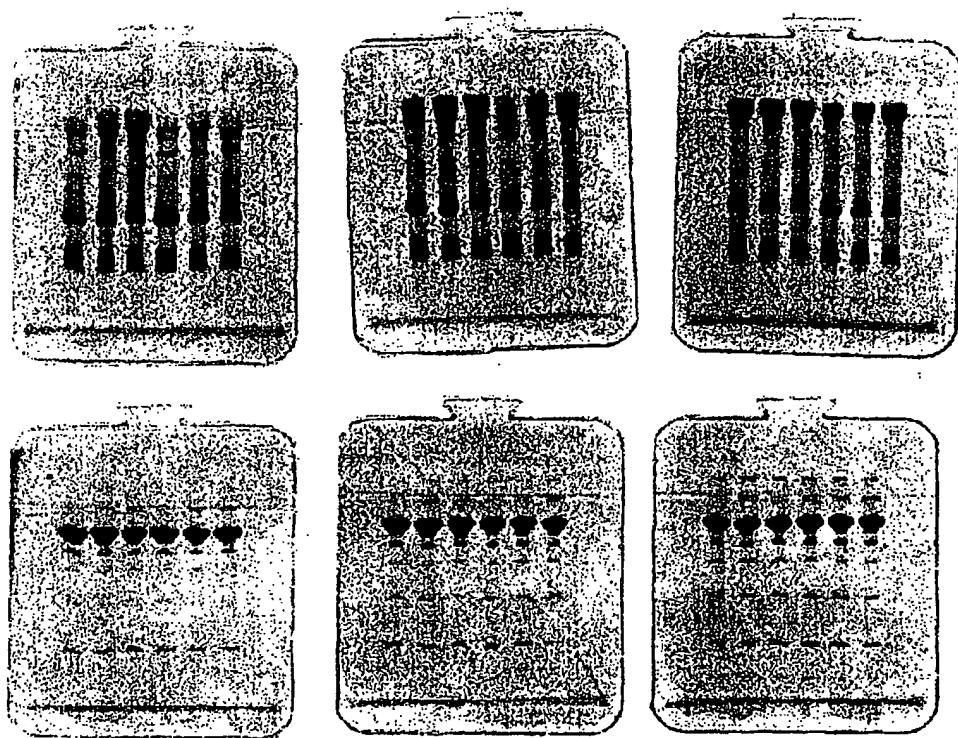


FIG. 2



## INTERNATIONAL SEARCH REPORT

International application No.

PCT/JP00/09339

## A. CLASSIFICATION OF SUBJECT MATTER

Int.Cl.<sup>7</sup> A61K39/395, 9/08, 47/04, 47/12

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

Int.Cl.<sup>7</sup> A61K39/395-39/44, 9/00-9/08, 47/00-47/28

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)  
CAPLUS (STN), MEDLINE (STN), EMBASE (STN), BIOSIS (STN),  
BIOTECHABS (STN), WPI (DIALOG)

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	JP, 4-228089, A (Kanegafuchi Chem. Ind. Co., Ltd.), 18 August, 1992 (18.08.92),	1, 6
Y	Claims; example (Family: none)	2-5, 7-14
Y	WO, 89/11298, A1 (CENTOOR, INC.), 30 November, 1989 (30.11.89), Claims; page 7, line 22 to page 8, line 2; example & JP, 3-504499, A Claims; page 3, lower left column, lines 16 to 23; example & EP, 417191, A	1-14
Y	WO, 89/11297, A1 (CENTOOR, INC.), 30 November, 1989 (30.11.89), Claims; example & JP, 3-504605, A Claims; example & EP, 417193, A	1-14
Y	WO, 96/22790, A1 (XENOTECH INCORPORATED), 01 August, 1996 (01.08.96), Claims; page 12, line 1 to page 17, line 27; page 28, line 14 to page 30, line 4	1-14

☒ Further documents are listed in the continuation of Box C. ☐ See patent family annex.

\* Special categories of cited documents:  
 "A" document defining the general state of the art which is not  
 considered to be of particular relevance  
 "E" earlier document but published on or after the international filing  
 date  
 "L" document which may throw doubts on priority claim(s) or which is  
 cited to establish the publication date of another citation or other  
 special reason (as specified)  
 "O" document referring to an oral disclosure, use, exhibition or other  
 means  
 "P" document published prior to the international filing date but later  
 than the priority date claimed

"T" later document published after the international filing date or  
 priority date and not in conflict with the application but cited to  
 understand the principle or theory underlying the invention  
 "X" document of particular relevance; the claimed invention cannot be  
 considered novel or cannot be considered to involve an inventive  
 step when the document is taken alone  
 "Y" document of particular relevance; the claimed invention cannot be  
 considered to involve an inventive step when the document is  
 combined with one or more other such documents, such  
 combination being obvious to a person skilled in the art  
 "&" document member of the same patent family

Date of the actual completion of the international search  
30 January, 2001 (30.01.01)Date of mailing of the international search report  
20 February, 2001 (20.02.01)Name and mailing address of the ISA/  
Japanese Patent Office

Authorized officer

Facsimile No.

Telephone No.



## INTERNATIONAL SEARCH REPORT

International application No.

PCT/JP00/09339

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
	& JP, 10-512885, A Claims; page 17, line 14 to page 22, line 1; page 31, line 5 to page 32, line 2 & AU, 9647645, A & US, 5626845, A & US, 5993817, A & EP, 813423, A1 & KR, 98701585, A	
Y	JP, 11-80025, A (Chugai Pharmaceutical Co., Ltd.), 23 March, 1999 (23.03.99), Claims; Par. Nos. [0002], [0062] & WO, 98/51329, & EP, 1004313, A1 & AU, 9872369, A & NO, 9905558, A	1-14
Y	WO, 00/00219, A1 (Chugai Pharmaceutical Co., Ltd.), 06 January, 2000 (06.01.00), Claims; page 2, lines 5 to 15; page 2, the last line to page 3, the last line; page 16, line 18 to page 17, line 21 & AU, 9942899, A	1-14
Y	WO, 98/13388, A1 (Chugai Pharmaceutical Co., Ltd.), 02 April, 1998 (02.04.98), Claims; page 1, line 18 to page 2, line 23; example & JP, 11-92500, A & EP, 962467, A1 & ZA, 9708590, A & AU, 9743972, A & NO, 9901449, A & CN, 1237983, A	1-14
Y	WO, 96/33735, A1 (CELL GENESYS, INC.), 31 October, 1996 (31.10.96), Claims; example 7 & JP, 11-505523, A Claims; example 7 & EP, 822830, A1 & AU, 9656322, A & KR, 99008096, A & US, 6075181, A	1-14
Y	WO, 96/39184, A1 (THE REGENTS OF THE UNIVERSITY OF CALIFORNIA), 12 December, 1996 (12.12.96), Claims; page 1, lines 17 to 26; page 11, lines 8 to 19 & US, 5660826, A & AU, 9658844, A	1-14
Y	SATO, K., et al., "Passive Immunization with Anti- Parathyroid Hormone-Related Protein Monoclonal Antibody Markedly Prolongs Survival Time of Hypercalcemic Nude Mice Bearing Transplanted Human PTHrP-Producing Tumors", J. Bone Miner. Res., 1993, 8 (7), pp. 849-860, ABSTRACT	1-14
PX	JP, 2000-80100, A (JAPAN TOBACCO INC.), 21 March, 2000 (21.03.00), Claims; Par. Nos. [0005], [0009], [0013], [0014], [0055], [0056]; example (Family: none)	1-8

Form PCT/ISA/210 (continuation of second sheet) (July 1992)